Impact of community-based vector control on house infestation and *Trypanosoma cruzi* infection in *Triatoma infestans*, dogs and cats in the Argentine Chaco


*Laboratorio de Eco-Epidemiología, Facultad de Ciencias Exactas y Naturales, Universidad de Buenos Aires (UBA), Buenos Aires, Argentina*

*b Instituto Nacional de Parasitología “Dr. Mario Fatala Chaben”, Paseo Colón 568, 1063 Buenos Aires, Argentina*

*c Division of Epidemiology and Preventive Medicine, Department of Veterinary Pathobiology, University of Illinois, Urbana-Champaign, 2001 South Lincoln Avenue, Urbana, IL 61802, USA*

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Abstract

The relative impact of two community-based vector control strategies on house infestation by *Triatoma infestans* and *Trypanosoma cruzi* infection in bugs, domestic dogs and cats was assessed in two neighboring rural areas comprising 40 small villages and 323 houses in one of the regions most endemic for Chagas disease in northern Argentina. The prevalence and abundance of domestic infestation were 1.5- and 6.5-fold higher, respectively, in the area under pulsed, non-supervised control actions operating under the guidelines of the National Vector Control Program (NCVP) than in the area under sustained, supervised surveillance carried out jointly by the UBA research team and NCVP. The prevalence of infestation and infection varied widely among village groups within each area. In the pulsed control area, the prevalence of infection in bugs, dogs and cats was two- to three-fold higher than in the area under sustained surveillance, most of the infected animals qualified as autochthonous cases, and evidence of recent transmission was observed. Infection was highly aggregated at the household level and fell close to the 80/20 rule. Using multiple logistic regression analysis clustered by household, infection in dogs was associated positively and significantly with variables reflecting local exposure to infected *T. infestans*, thus demonstrating weak performance of the vector surveillance system. For high-risk areas in the Gran Chaco region, interruption of vector-mediated domestic transmission of *T. cruzi* requires residual insecticide spraying that is more intense, of a higher quality and sustained in time, combined with community participation and environmental management measures.

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1. Introduction

Triatoma infestans, the main vector of Chagas disease in the southern cone of South America, has been the target of various vector control programs for several decades. The Southern Cone Initiative, a regional effort to eliminate *T. infestans* and to interrupt transmission of *Trypanosoma cruzi*, was launched in 1991 (Dias et al., 2002). At present, vector-mediated transmission to humans has been interrupted in Chile, Uruguay, Brazil and parts of Argentina and Paraguay (Scholfield et al., 2006; WHO, 2002). However, transmission was not interrupted in the Gran Chaco, a natural landscape unit extending over northern Argentina, Bolivia, Paraguay and southwestern Brazil, where four million people live in a subsistence economy (Gürtler, in press).

Chagas disease vector control programs have typically not achieved the desired levels of vector control or suppression mainly because of interruption in provision of funding and resources, and due to reduced political will (Dias et al., 2002). In Argentina, the National Vector Control Program (NVCP) of *T. infestans* was implemented with a vertical structure in the early 1960s (Segura, 2002). Following disorganized and incomplete decentralization in the early 1980s, the Argentine NVCP adopted in 1992 a new vector control strategy (“Plan Ramón Carrillo”). This program was based on training of local villagers and primary health care agents in insecticide spraying activities and in monitoring house reinfestation (Segura, 2002). Community leaders and primary health care agents were provided with manual compression sprayers and pyrethroid insecticides, and kept records of bug notifications and insecticide treatments over time. Vector control activities comprised an initial attack phase, in which all houses in a village were scheduled to be sprayed twice over a 6-month period, and a surveillance phase, in which all houses were scheduled for monitoring of reinfestation and selective sprays. Over, 1993–2001, all of the 961,500 rural houses in the endemic area were reported to have been sprayed with insecticides during the attack phase, and 85% of the houses were considered to be under surveillance (Segura, 2002). This community-based vector control program was gradually discontinued during the late 1990s, and the reported number of Chagas acute cases subsequently increased, most notably in Santiago del Estero and other neighboring provinces in the Argentine Chaco. Between 1998 and 2004, insecticide spraying rates have remained low at some 60,000 houses per year, and since then vector control actions have been rather erratic and mainly dependent on the political will, knowledge and budget of each province. The actual impact of cumulative control actions on domestic and peridomestic infestation with *T. infestans* and *T. cruzi* transmission has not been assessed.

As part of a longitudinal study aimed at modeling the transmission dynamics and control of *T. cruzi* in a well-defined rural area in Santiago del Estero, all houses from five neighboring villages (the core area, centered in Amamá village) have been regularly monitored for reinfestation over a decade after community-wide residual application of insecticides in 1992 (Gürtler et al., 1999; Cohen and Gürtler, 2001). The vector surveillance program was based on community participation and professional focal spraying in 1993–1995, and on selective insecticide treatments conducted by householders and community leaders between 1996 and 2004 (Cecere et al., 1999, 2002). The long-term impact of such actions on domestic and peridomestic infestation with *T. infestans* and *T. cruzi* transmission was assessed by means of entomological indices and by using domestic dogs as natural sentinels of transmission (Cardinal et al., 2006a) to avoid regular screening surveys of the local human populations. The prevalence of *T. cruzi* infection in dogs decreased from 65.1% at baseline to 4.7% at 10 years after sustained vector surveillance, whereas the average annual force of infection dropped 260-fold from 72.7 per 100 dog-years at baseline to <0.3% in 2002 (Gürtler et al., 2005; Cardinal et al., 2006a).

In this study, we assessed the relative impact of two community-based vector control strategies on infestation and transmission of *T. cruzi* to *T. infestans*, domestic dogs and cats in two neighboring rural areas. The core area was under sustained, community-based surveillance supervised jointly by the University of Buenos Aires (UBA) research team and NCVP since 1992. The peripheral area, operating under the guidelines of the “Plan Ramón Carrillo” program between 1993 and 1999 and variations of it thereafter, experienced pulsed, non-supervised, community-based insecticide applications promoted by NCVP. Because of the fast recovery rate of local *T. infestans* populations and of domestic transmission in the absence of sustained surveillance (Gürtler et al., 2005), we hypothesized that domestic infestation and transmission would be more intense in the area under pulsed, non-supervised vector control actions, and in those households having infected dogs or cats.

2. Materials and methods

2.1. Study area

Field studies were carried out in several villages around Amamá (27°12′33″S, 63°02′10″W), Moreno.
department, in the Province of Santiago del Estero, Argentina. The Moreno department reported the highest number of human acute cases of Chagas disease notified in Argentina. The study area has been described elsewhere (Ceballos et al., 2006; Cecere et al., 2002). Briefly, two areas were visited: (i) the core, including Amamá, Trinidad, Mercedes, Villa Matilde and Pampa Pozo, with 137 houses, and (ii) the periphery surrounding the core area, which includes 33 villages and 2 isolated settlements with 186 houses (La Celestina, San Ramón, San Isidro, El Rosario, San Salvador, Fracción El Rosario, Armonía, San Ignacio, San Pedro, Santos Lugares, San Roque, Santo Domingo, Gondra, Cañada Central, Central Dolores, Esperanza, La Dormida, Laprida, Las Cañitas, Villa Isabel, San Pablo, San Luis, Barrio Pampa Pozo, La Curva, Lote S, Dolores, Escudero, La Favorita, La Nueva Fortuna, Tacoyura, Santa Ana, Santa María, Santo Domingo de los Cisneros, Turquía and Villa Napenay). In the peripheral area, official vector control activities have been conducted mostly in two pulses (in 1993–1996 under “Plan Ramón Carrillo”, and in 2000–2001, when the recent occurrence of symptomatic acute cases of Chagas disease prompted a new round of residual spraying with insecticides. As part of the current investigation, in April 2004, all houses were sprayed with pyrethroid insecticides by NVCP personnel supervised by the research team and monitored for reinfestation thereafter.

2.2. Triatomine surveys

Two teams of four people each visited a total of 186 houses of the peripheral area in October–November 2002. Nearly all (93.4%) of the 137 houses from the core area had been surveyed in October and the results were reported (Cardinal et al., 2006a). A questionnaire inquiring on householders’ practices of vector control and perception of triatomine infestation was completed for each household as described elsewhere (Cecere et al., 1998). Briefly, villagers were asked when the last insecticide spraying of the house was done, who did it, which structures were sprayed, which insecticide was applied and the amount of insecticide used.

Skilled bug collectors from NVCP searched for triatomine bugs in all bedrooms (one person) and in peridomestic areas (two persons) of all compounds in which householders were present (n = 176) using a dislodgant spray (0.2% tetramethrin, Icona, Buenos Aires, Argentina) for 30 min per compound (Gürtler et al., 1999). Peridomestic structures included corrals for goats, sheep, cows, horses and pigs, chicken coops, trees where chicken roosted, storerooms, kitchens and other possible refuges for triatomines within the area of human activity. Householders were encouraged to collect invading triatomines, and were given plastic bags to keep them until our next visit. All bugs were later identified to species and stage at the field laboratory and counted per site. All live or moribund third to fifth instars and adults of T. infestans were individually examined for T. cruzi infection within 10 days of capture; Triatoma guasayana and T. gariabesi bugs were separately examined in pools of ≤3 insects from the same site. Bug feces were diluted with saline solution and microscopically examined at 220–400× magnification. Minicircle-DNA-based polymerase chain reaction applied to fecal lysates from T. infestans detected T. cruzi in 91% of microscope-positive bugs and only in 3.4% of microscope-negative bugs, but not in the other triatomine species collected (Marcet et al., 2006; Schijman et al., 2006).

2.3. Domestic animal surveys

A subset of the villages (58%) inspected for infestation and less distant from Amamá was chosen for the domestic animal surveys. A house-to-house census of all dogs and cats was undertaken in three surveys (November 2002, March and July 2003) totaling 103 houses from 17 villages and 2 isolated settlements of the periphery. A standard demographic questionnaire was completed for each animal (Cardinal et al., 2006a). Villagers were asked the name, age, sex and place of birth of each dog and cat they owned; the name of the animal’s mother, the animal’s history of visiting or residing in villages outside the study area, the animal’s main function in the household and other reproductive and behavioral details.

A total of 221 (85%) of 260 dogs and 61 (76%) of 80 cats was examined for infection. All animal and blood sample processing were conducted according to the Institutional Animal Care and Use Committee protocol no. 04223 at University of Illinois at Urbana-Champaign as reported elsewhere (Cardinal et al., 2006a). Thirty dogs and 18 cats aged <3 months were examined only by xenodiagnosis, whereas older animals were bled by venipuncture and diagnosed serologically; a subset of the latter was also examined by xenodiagnosis (124 dogs and 30 cats). Five dogs and 10 cats older than 3 months that could not be bled were examined only by xenodiagnosis. Six seropositive dogs were examined by xenodiagnosis in March or July 2003 to confirm T. cruzi infection and to isolate parasites; one seropositive dog died before being examined by xenodiagnosis.
2.4. Serodiagnosis and xenodiagnosis

Sera from dogs were tested for antibodies to *T. cruzi* by indirect hemagglutination assay (IHA; Polychaco, Buenos Aires, Argentina), indirect immunofluorescence test (IFAT) and enzyme-linked immunosorbent assay (ELISA) using standardized procedures and criteria reported elsewhere (Lauricella et al., 1998). Sera from cats were tested for antibodies to *T. cruzi* by IHA (Polychaco, Buenos Aires, Argentina), IFAT (total antigamma LID, Laboratorio Inmunodiagnóstico, Buenos Aires, Argentina) and ELISA (goat anti-cat IgG HRP, Santa Cruz Biotechnology, Santa Cruz, California). Titers ≥1:16 (IHA and IFAT) or an optical absorbance ≥0.2 (ELISA) were used as cut-off values for dogs and cats (Lauricella et al., 1998). Xenodiagnosis was performed using 20 laboratory-reared, third- or fourth-instar nymphs of *T. infestans* per animal for 25 min (Gürtler et al., 2007). Pools of feces from five bugs each that fed on a given animal were examined for *T. cruzi* infection at 400× magnification 30 and 60 days after feeding. Bugs from each positive pool were re-examined individually. ‘Seropositive’ refers to samples reactive by at least two different serologic tests among ELISA, IHA or IFAT. ‘Infected’ means animals having positive xenodiagnosis and/or being seropositive to *T. cruzi*. The results of serodiagnosis and xenodiagnosis were combined to calculate the composite prevalence of *T. cruzi* infection.

2.5. Data analysis

Fisher’s exact tests were used to compare proportions of age classes, dog-to-human ratios, infestation or infection between interventions areas, or proportions of infection between dogs and cats. Mann–Whitney tests were used to compare mean host numbers per house in infection between dogs and cats. Mann–Whitney tests were used to compare mean number of visits from the NVCP personnel and mean number of visits from the NVCP personnel for comparisons. Villages within each intervention area were grouped between intervention areas. Adjusted OR and CI were estimated from maximum-likelihood logistic multiple regression analysis clustered on dog’s compound to provide robust standard errors (Stata 9.0, College Station, TX). Backward and forward stepwise procedures were used to obtain the most parsimonious model that retained independent variables at the 5% nominal significance level. Interaction terms were then added to this model and tested for significance.

3. Results

A total of 1408 people was registered in the study area (Table 1). The mean number of people per house (4.8–4.6 persons) and the percentage younger than 15 years of age (40–45%) did not differ significantly between areas. House construction materials were similar between areas, and wall plasters were very cracked or nonexistent in 40–48% of the houses. The area under pulsed control actions had significantly more thatched
Table 1
Characteristics of two neighboring rural areas under different community-based vector control strategies in Santiago del Estero, October 2002

<table>
<thead>
<tr>
<th>Area under sustained surveillance</th>
<th>Area under pulsed control actions</th>
<th>Significance test</th>
</tr>
</thead>
<tbody>
<tr>
<td>Total population (% &lt;15 years)</td>
<td>595 (45%)</td>
<td>813 (40%)&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td>Mean number of people per house (S.D.)</td>
<td>4.8 (2.5)</td>
<td>4.6 (3.0)&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
<tr>
<td>Dog-to-human ratio (total number of dogs)</td>
<td>0.47 (279)</td>
<td>0.58 (475)&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td>Houses with wall plaster very cracked or nonexistent</td>
<td>40%</td>
<td>48%&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td>Houses with thatched roof</td>
<td>87%</td>
<td>91%&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td>Mean number of peridomestic sites per house (S.D.)</td>
<td>4.2 (2.1)</td>
<td>5.4 (3.0)&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
<tr>
<td>Mean number of chickens per house (S.D.)</td>
<td>17.6 (16.1)</td>
<td>20.2 (14.9)&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
<tr>
<td>Mean number of goats per house (S.D.)</td>
<td>6.2 (9.3)</td>
<td>8.2 (11.7)&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
<tr>
<td>Mean number of visits from NVCP personnel (S.D.)</td>
<td>9.4 (1.1)</td>
<td>3.5 (1.5)&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
</tbody>
</table>

<sup>a</sup> Mann–Whitney test.
<sup>b</sup> Fisher’s test.

roofs, peridomestic sites and chickens per house than the area under sustained surveillance. A higher ratio of dogs to humans and mean numbers of goats per house were also recorded in the area under pulsed control actions. According to NVCP records, no houses were sprayed in the pulsed control actions area during 1996–1998; spray coverage was low during 1999 (8%), peaked in 2000 (44%), decreased in 2001 (7%), and no spraying actions were recorded thereafter. However, in the area under sustained surveillance over the 5-year period 1997–2002, selective insecticide spraying of domiciles was always >0% and ranged from 6 to 27%. Villages under pulsed control actions experienced significantly less visits from NVCP personnel between 1993 and 2002 (3.5 visits) than villages under sustained surveillance (9.4 visits) (Table 1).

The prevalence and abundance of domestic infestation were 1.5- and 6.5-fold higher, respectively, in the area under pulsed control actions than in the area under sustained, supervised surveillance (Fig. 1A). Infestations in nearby peridomestic sites (\( P = 0.009 \)) and corrals (\( P = 0.004 \)) in the pulsed control area were significantly higher than in the area under sustained surveillance. Chicken coops and goat or pig corrals were the peridomestic structures most frequently found infested and with the highest bug abundance in each area.

The prevalence of *T. infestans* infected with *T. cruzi* in domiciles was 2.9-fold higher in the area under pulsed control actions (13.2%) than in the area under sustained surveillance (4.5%) (\( P = 0.04 \)) (Fig. 1B). Bug infection Fig. 1. (A), Prevalence and abundance of *Triatoma infestans* and (B), prevalence of *Trypanosoma cruzi* infection in *Triatoma infestans*, dogs and cats from two neighboring rural areas under different vector control strategies in Santiago del Estero, October 2002–July 2003. The mean abundance of bugs and (standard deviation) are shown beside squares (A). Numbers on top of bars are individuals infected/examined for infection (B).
Table 2
Prevalence of infestation by *Triatoma infestans* and infection with *Trypanosoma cruzi* in *Triatoma infestans*, dogs and cats from two neighboring rural areas under different community-based vector control strategies in Santiago del Estero, October 2002–July 2003; nd, no data available.

<table>
<thead>
<tr>
<th>Village group</th>
<th>Number of houses</th>
<th>Prevalence of infestation (%)</th>
<th>Prevalence of infection (number examined)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Domestic</td>
<td>Peridomestic</td>
<td>Bugs</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Dogs</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Cats</td>
</tr>
<tr>
<td><strong>Area under pulsed control actions</strong>&lt;sup&gt;b&lt;/sup&gt;</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Celestina</td>
<td>7</td>
<td>0</td>
<td>57</td>
</tr>
<tr>
<td>Central Dolores</td>
<td>48</td>
<td>9</td>
<td>57</td>
</tr>
<tr>
<td>Escudero</td>
<td>22</td>
<td>10</td>
<td>40</td>
</tr>
<tr>
<td>Gondra</td>
<td>6</td>
<td>0</td>
<td>20</td>
</tr>
<tr>
<td>Laprida</td>
<td>20</td>
<td>5</td>
<td>63</td>
</tr>
<tr>
<td>La Curva</td>
<td>6</td>
<td>83</td>
<td>100</td>
</tr>
<tr>
<td>Pampa Pozo Barrio</td>
<td>2</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Rosario</td>
<td>39</td>
<td>37</td>
<td>49</td>
</tr>
<tr>
<td>San Pablo</td>
<td>8</td>
<td>0</td>
<td>25</td>
</tr>
<tr>
<td>Santo Domingo</td>
<td>26</td>
<td>23</td>
<td>46</td>
</tr>
<tr>
<td><strong>Total</strong></td>
<td>186</td>
<td>17.6</td>
<td>50.7</td>
</tr>
<tr>
<td><strong>Mean (S.D.)</strong>&lt;sup&gt;a&lt;/sup&gt;</td>
<td>18.5 (25.7)</td>
<td>50.7 (22.1)</td>
<td>3.6 (4.4)</td>
</tr>
<tr>
<td><strong>Area under sustained surveillance</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Amamá</td>
<td>66</td>
<td>12</td>
<td>35</td>
</tr>
<tr>
<td>Mercedes</td>
<td>34</td>
<td>6</td>
<td>30</td>
</tr>
<tr>
<td>Pampa Pozo</td>
<td>9</td>
<td>0</td>
<td>33</td>
</tr>
<tr>
<td>Trinidad</td>
<td>28</td>
<td>23</td>
<td>23</td>
</tr>
<tr>
<td><strong>Total</strong></td>
<td>137</td>
<td>11.7</td>
<td>31.3</td>
</tr>
<tr>
<td><strong>Mean (S.D.)</strong>&lt;sup&gt;a&lt;/sup&gt;</td>
<td>9.0 (9.9)</td>
<td>19.3 (11.6)</td>
<td>0.7 (0.9)</td>
</tr>
</tbody>
</table>

<sup>a</sup> Only villages with ≥6 houses or with ≥10 *T. infestans* or dogs examined for infection were considered when calculating mean infestation or infection prevalence.

<sup>b</sup> Excludes two isolated settlements where no *T. infestans* were found (San Luis and Lote S). No infection was found in three dogs and one cat examined from the latter.

Prevalence was highest in domiciles in both areas, lower in nearby peridomestic sites in the area under pulsed control actions (2.5%), and rare in corrals (<1.4%). The bug infection prevalence for all peridomestic structures pooled together was 1.8% (*n* = 737) for *T. infestans*. No *T. garciabesi* (*n* = 151) or *T. guasayana* (*n* = 25) were found infected with *T. cruzi*. In the area under sustained surveillance, *T. cruzi* infection was only detected in 2.5% of *T. infestans* males (*n* = 120); no infection was found among third or fourth instars (*n* = 74), fifth instars (*n* = 113) and females (*n* = 82). In the pulsed control area, infection increased from 0% in third or fourth instars (*n* = 175) to 3.3% in fifth instars (*n* = 243), and to 5.6% in males (*n* = 288) and 8.3% in females (*n* = 168).

The composite prevalence of *T. cruzi* infection in the pulsed control area relative to the sustained surveillance area was 2.5-fold higher in all dogs (11.8% versus 4.7%, *P* = 0.006) and 3.1-fold higher in all cats (6.6% versus 2.1%, *P* = 0.38) (Fig. 1B). In the pulsed control area, dogs (11.8%) had a 1.8-fold higher prevalence of *T. cruzi* than cats (6.6%), but this difference was also not statistically significant (*P* = 0.35).

Within each intervention area, the prevalence of infested houses and of *T. cruzi*-infected bugs, dogs and cats varied widely among village groups (Table 2). However, the area under pulsed control actions showed the highest prevalence of infestation in domiciles (83%) and peridomestic sites (100%), and the highest prevalence of infection in bugs (12.7%), dogs (20.0%) and cats (16.7%).

Of 103 households surveyed for domestic animals, 94% owned at least one dog and 61% at least one cat. The mean number of dogs and cats per household was 2.6 (S.D. = 1.5) and 0.8 (S.D. = 0.8), respectively. Dogs were 3.3 times more abundant than cats (260 versus 80, respectively) and nearly all of them were mongrels. The sex ratio between males and females (M:F) was significantly biased toward males in dogs (M:F = 5.8, *χ²* = 128.4, degrees of freedom [df] = 1, *P* < 0.0001) but not in cats (M:F = 1.4, *χ²* = 2.5, df = 1, *P* > 0.1). Both populations were very young, with median ages of 2.0 years (first and third quartiles [Q1–Q3] = 1–5) for dogs and 1.8 years (Q1–Q3 = 0–4) for cats. Cats (75%) were born locally significantly more frequently than dogs (57%).
Fig. 2. Age-specific prevalence of *Trypanosoma cruzi* infection in native dogs and cats from the area under pulsed control actions in Santiago del Estero, November 2002–July 2003. Numbers close to data points represent the numbers of animals examined for infection. Figure excludes seven dogs and three cats of unknown age. (P = 0.004). Most dogs and cats were unrestrained; 40% of the dogs were used for hunting; 50% of the dogs and 68% of the cats were reported to sleep indoors.

The age-prevalence curves in native cats and dogs from the area under pulsed control actions were very similar up to 5 years of age (Fig. 2). When host age was divided in two classes (≤1 and >1 year), multiple logistic regression analysis showed no significant effects on host *T. cruzi* infection of host type (dog or cat) (OR = 1.4, CI = 0.5–4.3) but significant effects of host age (OR = 3.6, CI = 1.6–8.1) (Wald $\chi^2 = 13.4$, 2 df, P < 0.005). Among native dogs from the area under pulsed control actions, the age-prevalence curve rose steadily from 5 to 10% at ages ≤1 year to 27% at age 3 years, and fluctuated between 13% and 25% in older ages (Fig. 3). In contrast, for the sustained surveillance area, the age-prevalence curve of native dogs younger than 10 years of age fluctuated below 5% and rose steeply to 35% only among dogs aged ≥10 years—the dogs that were alive before the 1992 insecticide spraying campaign (Cardinal et al., 2006a). In the pulsed control area, the prevalence of *T. cruzi* in immigrant dogs (7.5%, n = 94) was half of that in native dogs (15.0%, n = 127) and peaked at age 2 (22%) and >6 (17–25%) years old (data not shown). Most of the infected animals (77%) in the pulsed control area were natives with permanent residence, and therefore qualified as autochthonous cases.

The 30 infected dogs and cats detected were in 16 (15.5%) households from nine villages. Ten of these houses (63%) harbored *T. cruzi*-infected *Triatoma infestans*, three harbored uninfected bugs, two had no bugs detected, and from one house no entomologic data were collected. Only a very small fraction of the households surveyed harbored infected animals (9% for *T. infestans*, 3% for cats, 16% for dogs) (Fig. 4), resulting in high household aggregation of *T. cruzi* infection (test of variance, $\chi^2 = 228.4$, df = 96, P < 0.0001). All infected cats were found cohabiting with infected dogs (P = 0.004). One household had three infected dogs and two infected cats.

Serologic titers of dogs and cats showed a clear-cut distinction between non-reactive and reactive sera, with nearly perfect concordance between ELISA and IFAT. A total of 191 sera (163 dogs and 28 cats) with ELISA absorbance values <0.15 and IFAT titers ≤1:8 were clearly negative, whereas 25 sera (22 dogs and 3 cats) with ELISA ≥0.25 and IFAT ≥1:64 were clearly...
positive. The only serologically discordant dog (ELISA <0.15, IFAT \(\geq 64\) and IHA \(\leq 1:8\)) was not examined by xenodiagnosis. Concordance between xenodiagnosis and serodiagnosis was 96% (119/124) in dogs and 100% (n = 30) in cats. Four dogs aged 2–12 years old were seropositive for \(T. cruzi\) and xenodiagnosis-negative. Only one dog, a 5-month-old native pup with permanent residence, was xenodiagnosis-positive and seronegative by the three tests. This dog most likely was an acute case of infection, as it cohabited with two dogs and one cat all infected with \(T. cruzi\) in a house with infected domestic bugs.

The relationship between \(T. cruzi\) infection in dogs and significant risk factors by univariate analysis is shown in Table 3. Ages were unknown for 7 dogs and no entomological data were available for 10 dogs. Based on a multiple logistic regression analysis of data clustered by house, \(T. cruzi\) infection was associated positively and significantly with age of the dog, sex (being a female), having permanent residence in the pulsed control area, the number of infected dogs or cats the animal cohabited with, the pooled abundance of \(T. infestans\) in domestic or nearby peridomestic sites at the dog’s compound, and the proportion of infected \(T. infestans\) at the dog’s compound (Wald \(\chi^2 = 54.7, 7\) df, \(P < 0.0001\)) (Table 3). Living in houses that had only been sprayed twice was not a significant risk factor in the regression analysis. In univariate analyses, infection in dogs was not significantly associated with having a hunting habit, type of spraying operator, insecticide spray coverage, years since last insecticide spray, abundance of \(T. infestans\) in corrals, or with the capture of \(T. guasayana\) or \(T. garcia-besi\) at the dog’s compound (data not shown). Among 33 dogs for which the mother’s name and serostatus was known, three infected dogs (37.5%) had seropositive mothers, whereas only two infected dogs (8%) had seronegative mothers.

In cats, \(T. cruzi\) infection was associated positively and significantly with cohabiting with at least one infected dog or cat (OR = 58.2, CI = 2.8–1194.2), the abundance of \(T. infestans\) in nearby peridomestic sites at the cat’s compound (OR = 46.0, CI = 3.6–587.2), and the occurrence of infected \(T. infestans\) at the cat’s compound (OR = 103.0, CI = 4.8–2233.4). None of the studied variables in cats were significant in multiple regression analysis.

4. Discussion

Our study shows that active transmission of \(T. cruzi\) occurred in villages under pulsed, non-supervised, community-based vector control actions operating under
NCVP guidelines for a decade, but not in the neighboring villages that were under sustained, supervised, community-based surveillance during the same period of time. The prevalence of *T. cruzi* infection in *T. infestans*, dogs and cats was two- to three-fold higher in the area under pulsed control actions. Domestic infestations were also more frequent and abundant in the pulsed control area. Weak performance of the vector surveillance system in this area, including lack of full coverage of insecticide spraying, absence of supervision and irregular insecticide supply, led to a rapid recovery of the bug population after spraying and to renewed parasite transmission within 2–3 years of domestic reinfection. This pattern has been shown for a single village of the core area in the late 1980s (Gürtler et al., 2005). In the peripheral area, vector control actions were highly heterogeneous among villages and were mostly performed in two pulses. A recent analysis of the Argentinean vector control program in Moreno department (where the study villages are located) found that control actions during the attack phase lacked full coverage of rural villages and that reinestation was highly clustered in space (Vazquez-Prokopec et al., unpublished results).

Age-prevalence curves reflected the history of major vector control actions, despite the fact that activities were not homogeneous and simultaneous across all the villages and houses studied. In native dogs from the area under pulsed control actions, the prevalence of *T. cruzi* rose steadily from 5 to 10% at ages ≤1 year (born in 2001–2002) to nearly 25% at age 3 years or more (mostly born after the first round of insecticide spraying in 1993–1996). This curve reflects the massive vector control actions conducted in most of the study villages in two pulses; the 2000–2001 control pulse was prompted by the occurrence of symptomatic acute cases of human Chagas disease. This rapid recovery of domestic transmission of *T. cruzi* may be explained by the combined effects of fast domestic reinestation, presence of infected dogs and cats resting indoors, their high incidence of infection and high infectiousness to bugs, and frequent blood meals on them (Cecere et al., 2006; Gürtler et al., 2007).

The area under pulsed control actions was more rural (included more peridomestic sites, goats, chickens and dogs) and slightly inferior housing quality than the core area under sustained, supervised surveillance. Whether these small differences, in addition to marked differences in insecticide spraying rates between areas, would modify domestic transmission remains unknown and requires further research. Lifestyle and animal management practices were very similar in the two intervention areas. Unlike the area under sustained surveillance where native dogs with permanent residence had a significantly lower prevalence of infection (5%) than immigrant dogs (11%) (Cardinal et al., 2006a), in the pulsed control area most of the infected animals were natives with permanent residence. This pattern is consistent with the lower risk of infection for dogs residing in or emigrating from the area under sustained surveillance in comparison to native dogs residing in or emigrating from the pulsed control area. The finding of a likely acute case of *T. cruzi* infection in a pup inhabiting an infested house with infected bugs, dogs and cats provides further evidence of vector-mediated transmission occurring locally at the time of the survey. Interestingly, a similar prevalence of *T. cruzi* infection in dogs (15.1%) was recorded in a rural area under irregular control actions located some 200 km away (but in the same ecoregion) in Chaco province (Diosque et al., 2004).

Infection in dogs was associated positively and significantly with variables reflecting local exposure to infected bugs (Table 3), which suggests that the most likely route of infection was vector-mediated within the dog’s compound. *T. infestans* was the only species found infected with *T. cruzi* even after 10 years of vector control actions in both areas. Vertical transmission may possibly account for some of the very young infected dogs. The occurrence of a xenodiagnosis-positive pup born to a seropositive, xenodiagnosis-positive female dog was most likely due to vertical transmission because the pup was only 3 days old, which is less than the latent period of vector-mediated infection. Cohabiting with at least one infected dog or cat was also significantly associated with *T. cruzi* infection in dogs and cats, which reflects household aggregation of infection. Cats were probably involved in the same vector-borne domestic transmission cycle as were dogs, as suggested by the association of infestation and cat infection and by the similar age-prevalence curves of native dogs and cats (Gürtler et al., 2007). Further evidence in support of this hypothesis was obtained by molecular typing of *T. cruzi* isolates from infected dogs, cats and domestic *T. infestans* of the study area; most of them shared sublineage TCIIe (Cardinal et al., unpublished results).

The evidence here provided gives further support to the use of dogs as natural sentinels of domestic and peridomestic transmission of *T. cruzi* (Gamboa, 1967; Gürtler et al., 1990; Castañera et al., 1998; Cardinal et al., 2006a). Several studies are showing that infected dogs constitute a major domestic reservoir of *T. cruzi* and a risk factor for domestic transmission throughout Latin America (Gürtler et al., 2005, 2007; Crisante et al., 2006; Estrada-Franco et al., 2006). Consequently, the regular screening of domestic dog populations for *T.*
cruzi infection can be used to identify houses or clusters with infected dogs and high risk of transmission. A highly sensitive and specific immunochromatographic dipstick test may be used for this purpose (Cardinal et al., 2006b) and for in situ implementation of appropriate strategies, such as insecticide-impregnated collaring (Reithinger et al., 2006), parasitidal drug treatment, culling, or selective spraying of the house.

Our study is the first to show household aggregation of T. cruzi infection in dogs and cats, so far only reported for humans (Mott et al., 1976). Host infection was concentrated in a small fraction of households (15.5%) and fell close to the 80/20 rule; this rule describes the disproportionate contribution to infection prevalence (80% or more) made by a small fraction (20% or less) of the households (Woolhouse et al., 1997). High aggregation of T. cruzi infection in dogs and cats also occurred in the area under sustained surveillance, where all the infected animals (12 dogs and 1 cat) were in 8.4% of houses. Household aggregation of T. cruzi infection is expected to increase the basic reproductive number (R0) of infection and the intensity of control actions needed to interrupt transmission. However, taking this heterogeneity into consideration may increase the effectiveness of vector control activities if effective targeting of the key households that concentrate infected reservoir hosts and transmission is achieved.

In the history of control actions against vector-mediated transmission of T. cruzi, there is a clear trend from vertically structured programs toward more horizontal strategies in which the community plays a crucial role during the surveillance phase (Bryan et al., 1994). In this study, we show that a community-based strategy can indeed be quite successful in reducing or controlling domestic T. cruzi transmission (not in eliminating T. infestans) provided it is accompanied by timely sustained support and external supervision is ensured in time. It is however, this temporal synchronicity that is so difficult to achieve in endemic areas where the rural population is sparsely distributed and access through dirt roads is limited, particularly during the rainy season. For highly endemic areas in the Gran Chaco region such as Santiago del Estero, interruption of vector-mediated domestic transmission of T. cruzi requires residual insecticide spraying that is more intense, of a higher quality, sustained in time, and combined with community participation and environmental management measures.

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