

# Oral Exposure to *Trypanosoma cruzi* Elicits a Systemic CD8<sup>+</sup> T Cell Response and Protection against Heterotopic Challenge<sup>∇</sup>

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***Trypanosoma cruzi* infects millions of people in Latin America and often leads to the development of Chagas disease. *T. cruzi* infection can be acquired at or near the bite site of the triatomine vector, but *per os* infection is also a well-documented mode of transmission, as evidenced by recent microepidemics of acute Chagas disease attributed to the consumption of parasite-contaminated foods and liquids. It would also be convenient to deliver vaccines for *T. cruzi* by the oral route, particularly live parasite vaccines intended for the immunization of reservoir hosts. For these reasons, we were interested in better understanding immunity to *T. cruzi* following oral infection or oral vaccination, knowing that the route of infection and site of antigen encounter can have substantial effects on the ensuing immune response. Here, we show that the route of infection does not alter the ability of *T. cruzi* to establish infection in muscle tissue nor does it impair the generation of a robust CD8<sup>+</sup> T cell response. Importantly, oral vaccination with attenuated parasites provides protection against wild-type (WT) *T. cruzi* challenge. These results strongly support the development of whole-organism-based vaccines targeting reservoir species as a means to alleviate the burden of Chagas disease in affected regions.**

Millions of people throughout Latin America are affected by Chagas disease. This condition is caused by persistent infection with the hemoflagellate protozoan parasite *Trypanosoma cruzi*, which sustains infection in mammalian hosts by replication of amastigotes in a cytoplasmic niche (61). Accounting for nearly 700,000 disability-adjusted life years (DALY) (39), Chagas disease is a prominent public health challenge, and current approaches to treatment and prevention are far from optimal (52). Metacyclic trypomastigotes in the feces of triatomine vectors are infective to a wide range of mammals, which primarily acquire *T. cruzi* through breaks in the skin, exposure to mucosal surfaces, or ingestion (61). Domestic spraying campaigns have had some degree of success in controlling transmission, most notably in Chile, Uruguay, and parts of Brazil (17). In this context, more recent attention has been given to outbreaks of *T. cruzi* infection acquired from food or drink tainted with *T. cruzi*-laden triatomine excreta (14, 28, 67). These incidences confirm that oral infection with *T. cruzi* is not only possible, but may be even a major route of *T. cruzi* infection in humans. The prominence of oral infection among reservoirs is supported by observations of opossums (50), raccoons (55), and dogs (51) ingesting triatomine bugs. It is not known how frequently humans acquire *T. cruzi* infection from contaminated food or ingestion of fomites. From an epidemiological perspective, it would be beneficial to distinguish people who acquired *T. cruzi* by an oral route from those who acquired it by other means of transmission, perhaps by a phenotypic signature of the T cell response. This aim necessitates

a better understanding of the host immune response to *T. cruzi* following exposure via the gastrointestinal (GI) tract.

Oral infection may lead to distinct parasitological and/or immunological outcomes compared to other routes of infection. GI mucosal tissue forms an interface between the organism and its environment and constitutes an immense surface area constantly in contact with potential pathogens and commensals. Accordingly, the immune system associated with mucosae has evolved a unique capacity to determine when an aggressive response is appropriate, balancing regulation and activation (56). Antigen (Ag) encounter at the GI mucosa or in gut-associated lymphoid tissue often results in tolerance, particularly for T cell responses, a process largely mediated by the cytokines transforming growth factor  $\beta$  (TGF $\beta$ ) and interleukin-10 (IL-10) (27, 41). We questioned whether parasite-specific CD8<sup>+</sup> T cell responses may develop differently during oral infection and explored the possibility that *T. cruzi* may exhibit a tropism when infecting by the GI tract different than that previously observed with systemic routes of infection (68). T cell populations found in the mucosal tissue along the GI tract have several characteristics that separate them from T cells in peripheral circulation. For example, homing to the GI mucosa is controlled by expression of distinct adhesion molecules. T cells primed by dendritic cells (DC) from Peyer's patches (PP) or mesenteric lymph nodes (LN) (mesLN) express specific integrins on their surface that confer the ability to home to gut tissue (18, 45). We also asked if the mucosal route of infection would bias responding CD8<sup>+</sup> T cells to accumulate in the intestines, possibly at the expense of a parasite-specific CD8<sup>+</sup> response in other peripheral tissue.

If a robust immune response is generated with oral *T. cruzi* infection, one would hypothesize that vaccination by this route could be effective. Although not currently available, it is a major goal to develop a vaccine that protects against *T. cruzi* infection, especially one that elicits T cell-based immunity (42,

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63). CD8<sup>+</sup> T cells, which respond to foreign Ag processed from the intracellular compartment and presented on molecules from the class I major histocompatibility complex (MHC), are essential for controlling infection by *T. cruzi* (34). In mice, a population of immunodominant CD8<sup>+</sup> T cells recognizes epitopes derived from the *T. cruzi* trans-sialidase gene family (35), and CD8<sup>+</sup> T cells are strongly implicated in experimental vaccine-induced protection against this parasite (25, 43, 44, 62, 65). An attainable, practical approach to reducing the burden of Chagas disease would be to implement a transmission-reducing vaccine targeting reservoirs of *T. cruzi*. In areas where the disease is endemic, dogs are an integral component of domestic transmission of *T. cruzi* (13, 20, 22), and mathematical models have suggested that removing infected dogs from homes in some regions could almost completely prevent vector transmission to humans (13). To that end, our lab is working to generate genetically attenuated strains of *T. cruzi* as vaccine candidates. As this vaccine would likely be administered to reservoir dogs by feeding, we carried out protection studies with mice following oral immunization with an attenuated strain of *T. cruzi*.

Here, we address several specific questions. (i) Does entry via the oral route alter the capacity of *T. cruzi* to establish infection systemically in muscle tissue? (ii) Does the immune system recognize and respond to *T. cruzi* as a localized gut infection? (iii) Can the phenotype of *T. cruzi*-specific T cells be used to distinguish the route of exposure to *T. cruzi* in an already-infected host? (iv) Can oral vaccination with attenuated parasites induce protective immunity to systemic challenge with *T. cruzi*?

## MATERIALS AND METHODS

**Mice, parasites, infections, and vaccinations.** C57BL/6 (Ly5.2<sup>+</sup>) (B6) mice were purchased from either The Jackson Laboratory or National Cancer Institute at Frederick (Frederick, MD). Mice were maintained at the University of Georgia animal facility in microisolator cages under specific pathogen-free conditions. Mice were infected with metacyclic forms of Brazil or CL strain *T. cruzi* by oral gavage (p.o.), intraperitoneal (i.p.) injection, or subcutaneous infection in the footpad (f.p.). Mice could be infected by *ad libitum* ingestion of food that had been contaminated with *T. cruzi* metacyclic trypomastigotes (data not shown); however, a gavage needle was used in the majority of experiments to maximize infection efficiency and allow consistent dosing between animals. As previously described (6), epimastigote cultures were maintained in liver infusion tryptose (LIT) medium, and metacyclic trypomastigotes were generated by stressing cultures with triatomine artificial urine (TAU) medium. Mice were given 50,000 to 500,000 metacyclic trypomastigotes. Alternatively, for vaccination/challenge experiments, tissue culture trypomastigotes were obtained from passage through Vero cells, and mice were infected by i.p. or f.p. injection with 1,000 tissue culture trypomastigotes (TCT). Attenuated parasites (ECH1<sup>+/-</sup> ECH2<sup>-/-</sup>) were generated by targeting the tandem enoyl-coenzyme A (CoA) hydratase 1 and 2 genes in a high-throughput knockout system recently developed by our lab (66). These ECH1<sup>+/-</sup> ECH2<sup>-/-</sup> parasites fail to establish persistent infection in C57BL/6J mice (as determined by immunosuppression) (7) or in immunodeficient mice. For vaccine protection studies, attenuated ECH1<sup>+/-</sup> ECH2<sup>-/-</sup> metacyclic trypomastigotes were administered by gavage. Mice received 3 doses separated by approximately 2 weeks. A total of 5 × 10<sup>5</sup> metacyclic trypomastigotes were given for doses 1 and 3, and 1.35 × 10<sup>5</sup> metacyclic trypomastigotes were given in dose 2. Control mice received equal volumes (100 μl) of phosphate-buffered saline (PBS) at each dose. Mice were sacrificed by CO<sub>2</sub> inhalation, and cervical dislocation followed. All animal protocols were approved by the University of Georgia Institutional Animal Care and Use Committee.

**Assessing protection with *in vivo* imaging.** For protection studies, *T. cruzi* parasites of the CL strain expressing the far-red tdTomato protein (9, 64) were subcutaneously inoculated into superficial subcutaneous tissue of the footpads. Mouse feet were imaged every other day using the Maestro 2 *in vivo* imaging

system (CRI, MA) with the green set of filters (acquisition settings: 560 to 750 in 10-nm steps; exposure time of 88.18 ms and 2 × 2 binning). The total fluorescent signal was quantitated and normalized by exposure time and the area of the camera field corresponding to the source of the fluorescence, and values are reported as photons/cm<sup>2</sup>/second.

**Isolation of lymphocytes from nonlymphoid tissues and adoptive transfers.** Before tissue removal, mice were perfused with 20 ml of PBS containing 0.8% sodium citrate as an anticoagulant. Perfusion was done by opening the abdominal and thoracic cavities, nicking the portal vein, and forcing PBS into the heart ventricles with a one-half-inch 35-gauge needle and 10-ml syringe. Tissue-derived lymphocytes were obtained by teasing tissues apart and vigorously pushing them through a 40-μm nylon mesh screen. Lymphocytes were obtained from lamina propria and intestinal epithelium as previously described (33). In brief, small intestines were isolated, cleaned, and cut into small segments. Pieces were stirred at 37°C for 20 min in Ca<sup>+</sup>/Mg<sup>+</sup>-free Hanks balanced salt solution containing 10% fetal calf serum (FCS) and 1 mM dithioerythritol to free intraepithelial lymphocytes (IEL). Gut pieces were further digested by stirring for 1 h at 37°C in RPMI 1640 medium containing 5% FCS, 1 mM CaCl<sub>2</sub>, 1 mM MgCl<sub>2</sub>, and 150 units/ml type II collagenase (Sigma). Tissue homogenate was then passed over a 40-μm nylon mesh screen, and both IEL and lamina propria (LP) cell populations were further purified by collection from the interface of 44% Percoll in RPMI medium underlain with 67% Percoll in PBS.

**T cell phenotyping.** Spleens were homogenized, and red blood cells (RBCs) were lysed in a hypotonic ammonium chloride solution. Washes and staining were done in PAB (2% bovine serum albumin [BSA] and 0.02% azide in PBS). Peripheral blood was obtained by retro-orbital venipuncture or by nicking the tail and collecting blood with a capillary, collected in sodium citrate solution, and washed in PAB. Cells were obtained from peripheral tissue as described below. Cells were incubated with class I MHC tetramer-phycoerythrin (PE) complexes loaded with TSKb20 peptide (35) and the labeled antibodies (Abs). Cells were stained for 30 min at 4°C in the dark, washed in PAB, and fixed in 2% formaldehyde. For whole blood, RBCs were lysed after surface staining in a hypotonic ammonium chloride solution and washed twice in PAB. Data were acquired using a CyAn flow cytometer (DakoCytometry) and analyzed with FlowJo software (Tree Star). MHC I tetramer TSKb20 (ANYKFTLV)/K<sup>b</sup> was synthesized at the Tetramer Core Facility (Emory University, Atlanta, GA). Antibodies for flow cytometric analysis were purchased from BD Biosciences (San Jose, CA), eBioscience (San Diego, CA), and Caltag Laboratories/Invitrogen (Carlsbad, CA).

**PCR.** Quantitative real-time PCR was performed as previously described (16). Briefly, tissue was collected from mice and finely minced. Samples were incubated at 55°C for 4 h in SDS-proteinase K lysis buffer. DNA was extracted twice with phenol-chloroform-isoamyl alcohol (25:24:1), precipitated with 100% ethanol, and resuspended in nuclease-free water. Samples were analyzed on an iCycler (Bio-Rad). For real-time and standard PCR, the following primers were used to amplify a 182-bp product from genomic *T. cruzi* DNA: TCZ-F\*, 5'-GCTCTTGCCACAMGGGTGC-3', where M = A or C, and TCZ-R 5'-CCAAGCAGCGATAGTTCAGG-3' (16).

**Histology.** Heart and skeletal muscle was obtained from *T. cruzi*-infected mice and controls, fixed in 10% buffered formalin, and embedded in paraffin. Five-micrometer-thick sections were obtained and stained with hematoxylin-eosin (H&E). Inflammation was evaluated qualitatively according to the presence or absence of myocyte necrosis and the severity of leukocyte infiltration.

**Statistical analysis.** We calculated statistical significance with a two-tailed Student *t* test in all cases, and a one-way, nonparametric analysis of variance (ANOVA) was used for data shown in Fig. 6C.

## RESULTS

***T. cruzi* infects skeletal muscle following oral exposure.** We first asked if parasites infecting via the oral route establish a systemic infection similar to that generated using other routes of infection. Metacyclic trypomastigotes were administered to mice by oral gavage (p.o.). Control infections were done by injecting metacyclics intraperitoneally (i.p.) or subcutaneously in the footpad (f.p.). PCR analysis revealed parasite DNA in the skeletal muscle at 21 to 25 days postinfection (dpi) in both p.o. and f.p.-infected mice (Fig. 1A). In addition, similar patchy mononuclear cell infiltrates were detectable in skeletal muscle at 35 dpi in mice infected by p.o. and f.p. routes (Fig.

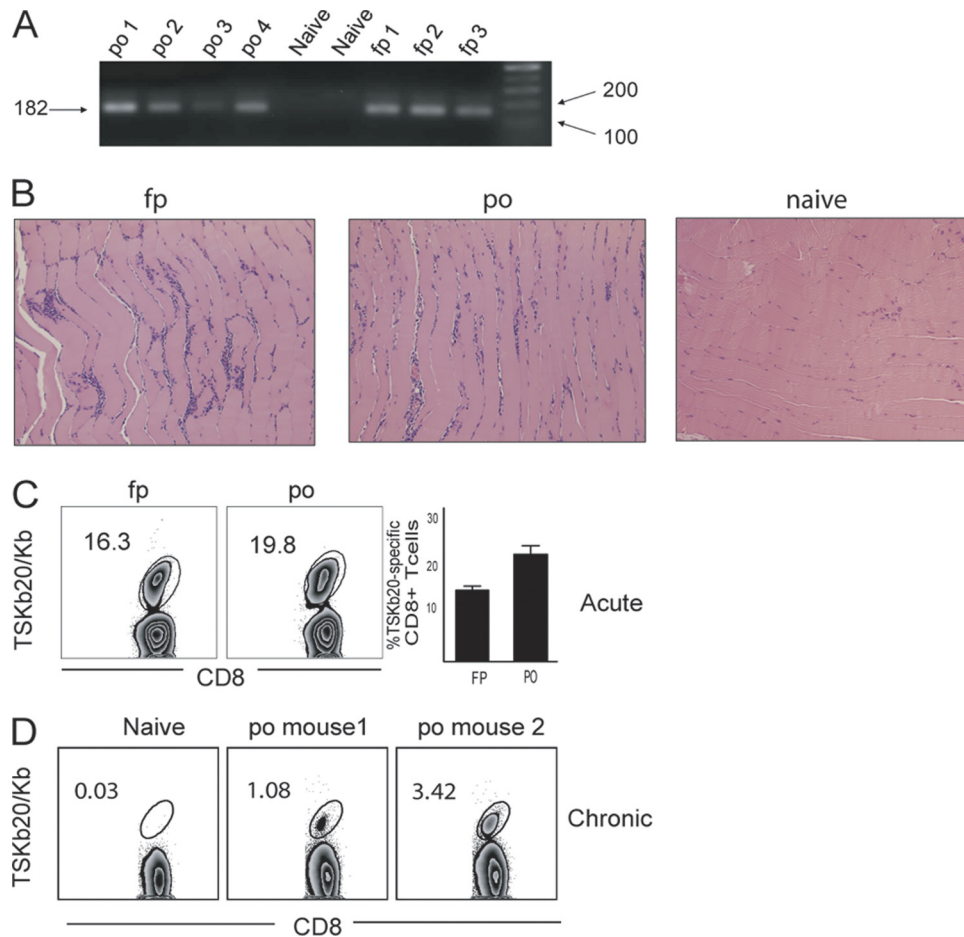


FIG. 1. Oral infection with metacyclic trypomastigotes of *T. cruzi* leads to systemic infection. (A) Metacyclic trypomastigotes delivered by either p.o. or f.p. route are detectable in skeletal muscle. DNA was extracted from skeletal muscle of mice 21 to 25 dpi, and a 182-bp segment of satellite DNA was amplified by PCR.  $n = 10$  for f.p. and  $n = 7$  for p.o. The sizes of the PCR product and markers are shown (in base pairs) at the left and right, respectively. (B) Similar inflammatory infiltrates develop in skeletal muscle of mice infected by f.p. and p.o. H&E sections of skeletal muscle are shown for naïve and p.o. and f.p.-infected mice at 35 dpi and are representative of at least 3 more experiments that ended between 21 and 40 dpi. (C and D) *T. cruzi* infection via the oral route induces a robust and lasting systemic T cell response. Representative flow plots show the frequency of CD8<sup>+</sup> T cells specific for TSKb20 in PBMC at 14 dpi ( $n = 10$  for p.o. and  $n = 5$  for f.p. [C]), and the means  $\pm$  standard error of the mean (SEM) are displayed. (D) TSKb20-specific CD8<sup>+</sup> T cells were measured in spleens of two individual mice infected p.o. at 140 dpi.

1B). The emergence of a population of CD8<sup>+</sup> T cells specific for the *T. cruzi*-derived immunodominant peptide TSKb20 (35) was also observed with mice infected p.o., comparable to mice infected in the f.p. (Fig. 1C). By 140 dpi, this robust TSKb20-specific CD8<sup>+</sup> T cell response contracted and stabilized at frequencies typical of chronic *T. cruzi* infection in mice (Fig. 1D) (35). Thus, within ~3 weeks after oral infection, *T. cruzi* disseminates, colonizes skeletal muscle, and induces a CD8<sup>+</sup> T cell response that is indistinguishable from that induced in mice infected via i.p. or f.p. routes.

***T. cruzi*-specific CD8<sup>+</sup> T cell traffic to tissues irrespective of infection route.** We hypothesized that parasite-specific CD8<sup>+</sup> T cells would accumulate in gut tissue following p.o. but not i.p. infection with *T. cruzi* if parasite Ag accumulated or persisted at the infection site. To examine this possibility, lymphocytes isolated from lymphoid and nonlymphoid tissues of mice infected by these routes were analyzed. The frequency and distribution of TSKb20<sup>+</sup> CD8<sup>+</sup> T cells did not vary depending on

infection route, as mice infected i.p. or by the f.p. developed parasite-specific CD8<sup>+</sup> T cells able to traffic to gut tissue, and mice infected p.o. had proportions of TSKb20-specific cells in spleen, lung, and skeletal muscle comparable to those in mice infected i.p. (Fig. 2A). We have recently shown that *T. cruzi*-specific CD8<sup>+</sup> T cells express CD69 selectively in sites of parasite persistence (M. H. Collins and R. L. Tarleton, unpublished data). CD8<sup>+</sup> T cells isolated from skeletal muscle of p.o., i.p., and f.p.-infected mice expressed similar levels of this marker of recent activation, whereas little expression was detected with T cells isolated from the spleens of these animals (Fig. 2B). Taken together, these results indicate that regardless of infection route, *T. cruzi* achieves a systemic infection and thus induces T cells capable of accessing multiple tissue compartments.

**T cells primed during oral *T. cruzi* infection are not imprinted for gut homing.** T cells are thought to be educated to adopt a particular homing pattern when Ag is acquired in



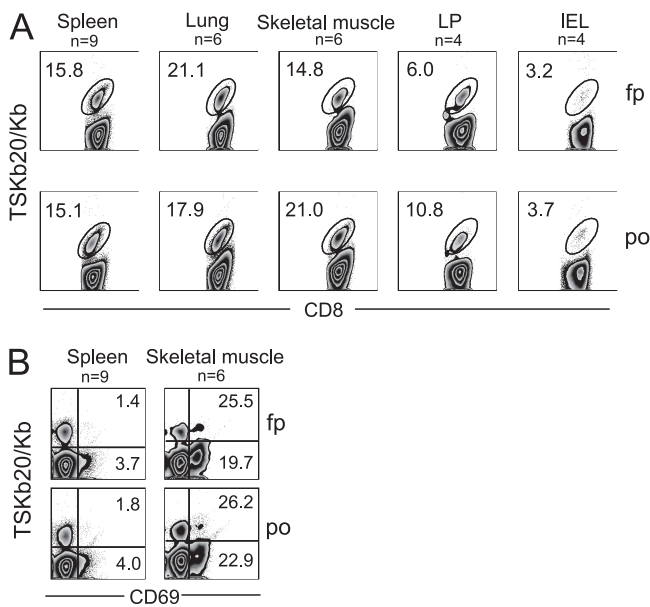


FIG. 2. *T. cruzi*-specific CD8<sup>+</sup> T cells traffic to peripheral tissues and are activated in sites of parasite persistence independently of infection route. (A) Flow plots show the frequency of TSKb20-specific CD8<sup>+</sup> cells in the indicated tissue of f.p.- or p.o. infected mice during acute *T. cruzi* infection. (B) Recently activated *T. cruzi*-specific CD8<sup>+</sup> T cells are present in sites of persistence. Flow plots are gated on CD8<sup>+</sup> T cells isolated from spleen or skeletal muscle of f.p.- or p.o. infected mice and show CD69 expression on TSKb20-specific cells. Numbers indicate the frequency of TSKb20<sup>+</sup> CD8<sup>+</sup> (top right) or TSKb20<sup>-</sup> CD8<sup>+</sup> (bottom right) T cells expressing CD69. Numbers in flow plots are the average frequency obtained from mice sacrificed between 19 and 39 dpi. The number of data points (*n*) is shown for each group; some data points comprise cells isolated from the pooled tissue of two or three mice harvested on the same day.

specific anatomical sites. Most notably, dendritic cells from GI tissue or GI-associated LN have been shown to confer a gut-homing program on T cells by induction of homing receptors such as CCR9 and  $\alpha 4\beta 7$  integrin (18, 45). To determine whether imprinting is a phenomenon associated with the initial T cell priming following oral exposure to *T. cruzi*, the expression of  $\alpha 4\beta 7$  was examined on parasite-specific CD8<sup>+</sup> T cells following p.o. and f.p. infection. Expression of this homing molecule was similar among TSKb20<sup>+</sup> CD8<sup>+</sup> T cells from spleen and mesLN in mice infected p.o. and by the f.p. with *T. cruzi* (Fig. 3B and C), suggesting that p.o. infection did not evoke a mucosally targeted CD8<sup>+</sup> T cell response.

While expression of  $\beta 7$  integrin is key for CD8<sup>+</sup> T cell homing to the LP and IEL compartments of the gut (30, 37), expression of CD103 ( $\alpha_E$  integrin) overlaps with that of  $\beta 7$  on Ag-specific CD8<sup>+</sup> T cells in the gut (38) and has been implicated in the localization of T cells in the intestine (59). CD103 was not expressed on TSKb20<sup>+</sup> CD8<sup>+</sup> T cells in peripheral blood at 14 dpi (Fig. 4A). Many (50% and 95%, respectively) CD8<sup>+</sup> T cells in LP and IEL populations are CD103<sup>+</sup>, but few TSKb20-specific CD8<sup>+</sup> T cells express this integrin (Fig. 4B)—though the frequency of tetramer-positive cells is likely diluted by CD8 $\alpha\alpha$ <sup>+</sup> T cells, especially in the IEL compartment (37). Importantly, the frequency of CD103-expressing cells is not affected by the route of infection. CD103 was also undetectable

on TSKb20<sup>+</sup> CD8<sup>+</sup> T cells in LN (data not shown). Expression of CD103 among total CD8<sup>+</sup> T cells was not enhanced in either peripheral (pLN) or gut-draining LN (mesLN) by an oral route of infection (Fig. 4C). Thus, no evidence of imprinting is observed after p.o. infection with *T. cruzi*.

Considering that CD69 is a marker of recent activation and that it associates with sphingosine-1-phosphate receptor 1 (S1P1) to retain lymphocytes in lymph nodes (60), we compared CD69 expression on CD8<sup>+</sup> T cells in pLN and mesLN as an alternative method for investigating anatomically restricted antigen (Ag) presentation. The expression of CD69 on TSKb20-specific CD8<sup>+</sup> T cells in pLN or mesLN was not favored by one infection route over the other (Fig. 4D). Taken together, all these data suggest that *T. cruzi* enters its host and disseminates early in infection; it is not anatomically confined or immunologically recognized as an infection of any one tissue compartment.

**An orally administered attenuated strain of *T. cruzi* provides protection against a challenge infection.** Given the robust CD8<sup>+</sup> T cell response observed with spleen, blood, gut, and other nonlymphoid tissues following p.o. infection by *T. cruzi*, we hypothesized that oral vaccination could provide protection against a heterologous route of *T. cruzi* challenge. To test this hypothesis, we exposed mice to an attenuated strain of *T. cruzi* by the oral route and subsequently challenged them with virulent parasites. Mice received three doses of *T. cruzi* strain CL attenuated by the deletion of one allele of the enoyl-CoA-hydratase-1 (ECH1) gene and both alleles of the ECH2 gene (66). Both ECH genes encode an enzyme involved in fatty acid oxidation, a process thought to be important in amastigote energy metabolism (2). Parasites deficient in ECH have been shown to exhibit poor replication as intracellular amastigotes *in vitro* and fail to establish persistent infection in most immunocompetent mice (D. Xu and R. L. Tarleton, unpublished data). Mice received ECH1<sup>+/-</sup> ECH2<sup>-/-</sup> parasites by oral gavage (oral vaccine) or control gavage solution (PBS). Two weeks after the final vaccination, peripheral blood mononuclear cells (PBMCs) from the majority of mice exhibited up-regulation of CD44, with over 50% of CD8<sup>+</sup> T cells becoming CD44<sup>hi</sup> in some cases (Fig. 5A). Vaccinated mice could be divided into two groups based on the presence of antigen-experienced CD8<sup>+</sup> T cells (with a 35% frequency of CD44<sup>+</sup> cells as the cutoff). CD8<sup>+</sup> T cells from “high responder” (HR) mice (*n* = 7) expressed this activation marker at significantly greater frequencies than “low responder” (LR) mice (*n* = 3) (47.1 versus 29.7, *P* = 0.029). Additionally, TSKb20<sup>+</sup> CD8<sup>+</sup> populations were apparent in the blood taken from one of the HR mice (Fig. 5B), further indicating the induction of *T. cruzi*-specific T cell responses in these mice. In contrast, LR mice were difficult to distinguish from PBS controls based on the CD44 expression profile of CD8<sup>+</sup> T cells (Fig. 5A) and did not have measurable levels of TSKb20-specific CD8<sup>+</sup> T cells in the peripheral blood.

The orally vaccinated mice were challenged in each footpad with  $2.5 \times 10^5$  CL trypomastigotes transfected with a gene encoding TdTomato fluorescent protein, and parasite load at the site of infection was monitored from 2 to 10 days postchallenge (dpc) by *in vivo* imaging (9). Orally vaccinated mice displayed a significant reduction in the fluorescence signal compared to control unvaccinated mice at all times measured

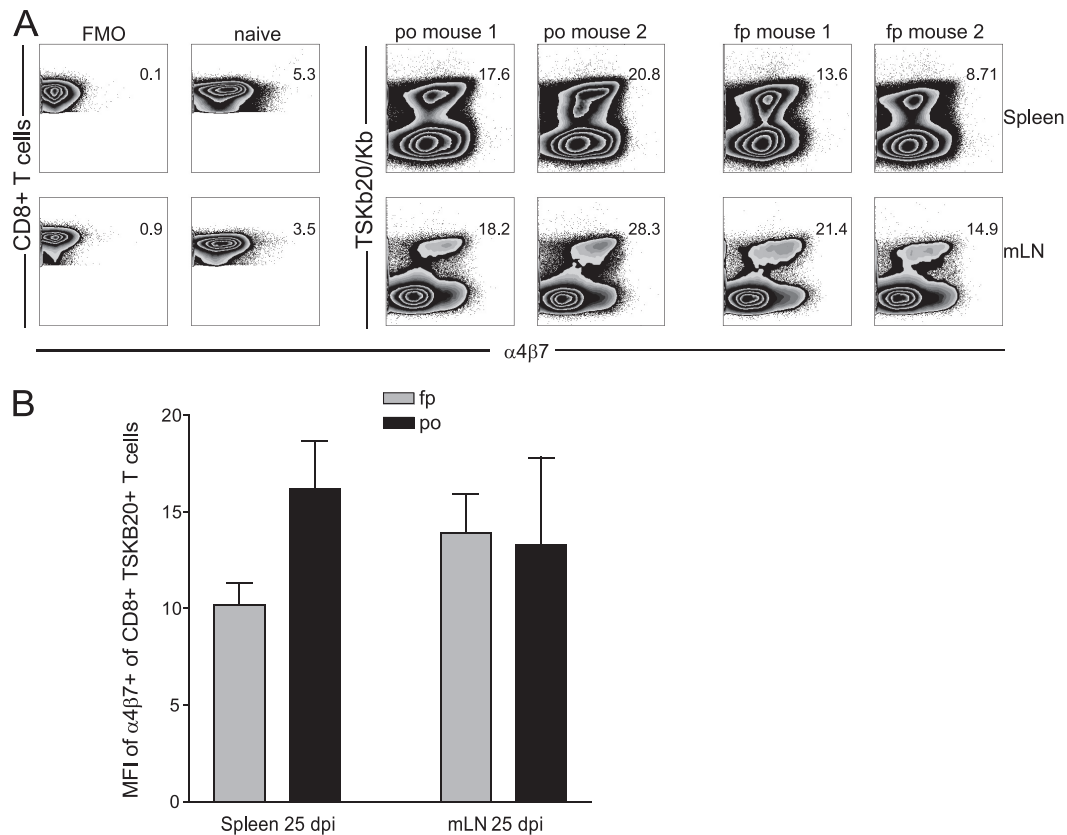


FIG. 3. Oral infection route does not skew *T. cruzi*-specific CD8<sup>+</sup> T cells toward a gut homing phenotype. Lymphocytes were isolated from spleen (top) and mLN (bottom) of *T. cruzi*-infected mice at 25 dpi and naïve age-matched mice, and surface expression of  $\alpha4\beta7$  was assessed by flow cytometry. (A) Fluorescence minus one (FMO) and naïve CD8<sup>+</sup> T staining controls for  $\alpha4\beta7$  expression are shown (left). Expression of  $\alpha4\beta7$  among TSKb20-specific CD8<sup>+</sup> T cells is shown for 2 representative mice from a total of 5 mice (right). Flow plots are gated on CD8<sup>+</sup> T cells. Numbers indicate the median fluorescence intensity (MFI) of CD8<sup>+</sup> T cells for FMO and naïve samples and the MFI of TSKB20<sup>+</sup> CD8<sup>+</sup> T cells for the orally or footpad-infected mice. (B) Mean  $\pm$  SEM of MFI for  $\alpha4\beta7$  expression on TSKB20<sup>+</sup> CD8<sup>+</sup> T cells. *n* = 5 mice per group.

(Fig. 6A and B). While the fluorescence signal progressively increased in PBS mice until 8 dpc, a consistently low signal was displayed from 2 to 6 dpc in oral vaccine mice, and measurements were statistically indistinguishable from those for the background (naïve) signal by 8 dpc. As expected, HR mice controlled f.p. infection with *T. cruzi* more efficiently than LR mice, but LR mice still displayed a significantly lower fluorescence signal than PBS mice at this time (Fig. 6C). At 25 dpc, oral vaccine mice exhibited lower parasite loads in skeletal muscle than the controls did, although both groups controlled parasites well after these early time points (Fig. 6D). Together, these results indicate that oral vaccination with attenuated parasites stimulated effective adaptive immunity that successfully protected mice from challenge with WT *T. cruzi*.

**DISCUSSION**

Mucosal surfaces are the target of infection for many human pathogens (46), including those that cause HIV, tuberculosis, respiratory tract infections, and diarrheal illnesses, and account for an enormous portion of global disease burden (39). Immune responses to infection or immunization may evolve differently relative to whether antigen exposure is initially or exclusively at mucosal or nonmucosal sites. For example, the

distribution of rotavirus and the responding cytotoxic T lymphocytes (CTL) varies according to route of infection (29, 47), and CD8<sup>+</sup>-dependent protection against mucosal HIV challenge can be generated by mucosal vaccination, whereas subcutaneous vaccination failed to elicit protective CTL at mucosal sites (5). Even simultaneously within the same animal, CD8<sup>+</sup> T cells specific for the same tumor Ag can acquire different homing phenotypes if the Ag is introduced in multiple distinct environments (8).

Oral *T. cruzi* infection in humans is an increasingly prominent public health issue (14, 28, 67), and the oral route may be the primary means of infection in many animal species. Unlike most strictly mucosal pathogens that remain localized to the airways, gut lumen, mucosal surface epithelium, or lamina propria, *T. cruzi* transits these superficial compartments and establishes a systemic infection. These characteristics of *T. cruzi* and the data from other infections predict that the immune system may respond differently to *T. cruzi* infection initiated orally compared to f.p. or i.p. infection. In this study, we investigated whether the CD8<sup>+</sup> T cell response to *T. cruzi* is altered by an oral route of parasite entry.

The nature of the immune response to parasitic infection can be heavily influenced by the earliest events of the host-parasite encounter, and there have recently been exciting ad-

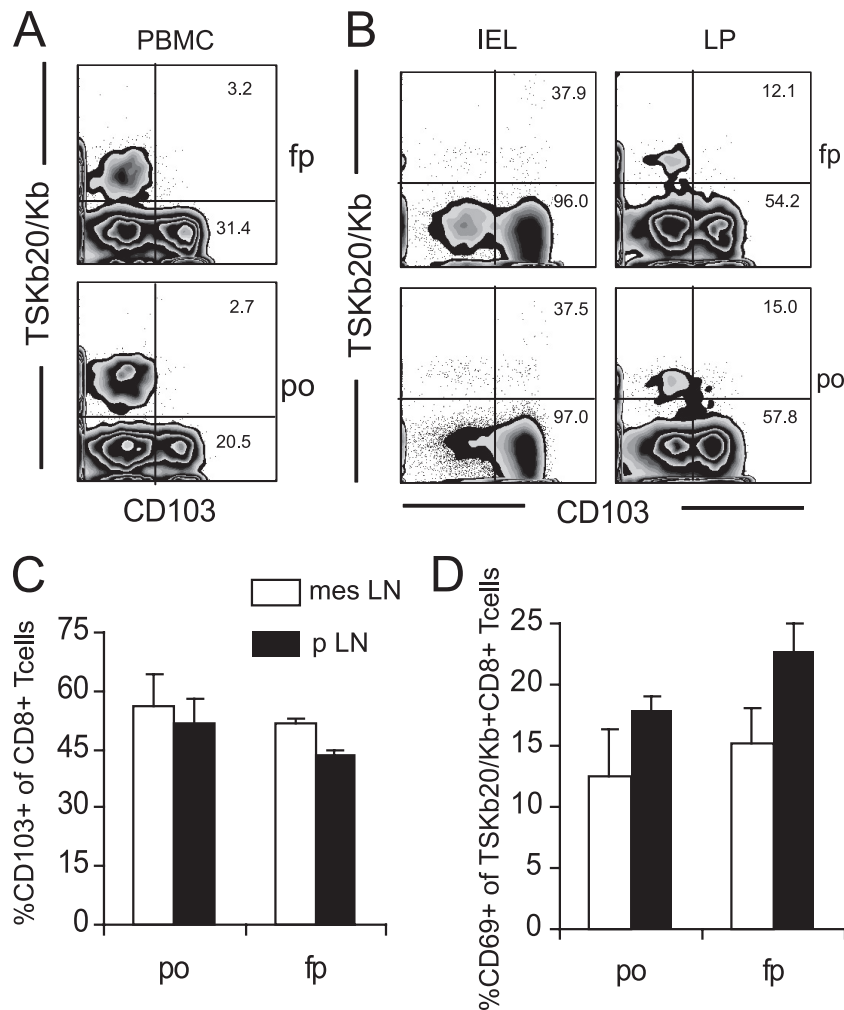
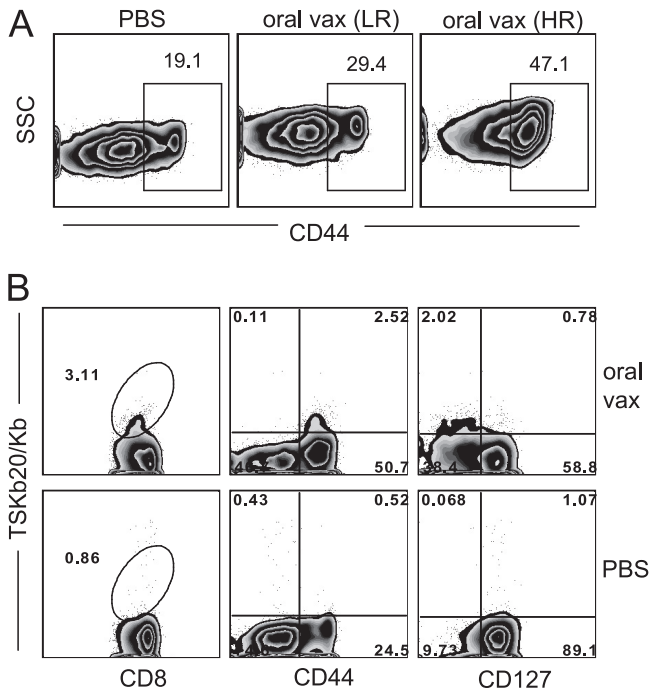


FIG. 4. Route of infection does not affect the distribution of TSKb20<sup>+</sup> CD103<sup>+</sup> CD8<sup>+</sup> T cells or activation of TSKb20<sup>+</sup> CD8<sup>+</sup> T cells in gut-draining LN. (A) Gut-imprinted CD8<sup>+</sup> T cells specific for *T. cruzi* are not detectable in blood early in infection. PBMC from mice at 14 dpi were isolated, and CD103 expression among TSKb20-specific cells was assessed by flow cytometry. Numbers in flow plots indicate the average percentage of TSKb20<sup>+</sup> CD8<sup>+</sup> (top) or TSKb20<sup>-</sup> CD8<sup>+</sup> (bottom) CD8<sup>+</sup> T cells that express CD103 (f.p.,  $n = 5$ ; p.o.,  $n = 10$ ). (B) CD103 is expressed by gut CD8<sup>+</sup> T cells, but by few *T. cruzi*-specific CD8<sup>+</sup> T cells. IEL and LP lymphocytes were isolated from *T. cruzi*-infected mice between 19 and 39 dpi, and surface expression of CD103 was assessed by flow cytometry. Flow plots are gated on CD8<sup>+</sup> T cells. Numbers indicate the average percentage of TSKb20<sup>+</sup> CD8<sup>+</sup> (top) or TSKb20<sup>-</sup> CD8<sup>+</sup> (bottom) CD8<sup>+</sup> T cells that express CD103, from 4 similar experiments. (C) Flow cytometry was used to assess the proportion of all CD8<sup>+</sup> T cells expressing CD103 in pLN and mesLN of mice after f.p. and p.o. infection with *T. cruzi*. Bars show average percentage of CD103<sup>+</sup> CD8<sup>+</sup> cells  $\pm$  SEM. (D) The proportion of recently activated TSKb20<sup>+</sup> CD8<sup>+</sup> T cells in different lymph nodes was assessed by flow cytometry. Bars show average percentage of CD103<sup>+</sup> CD8<sup>+</sup> cells  $\pm$  SEM.  $n = 3$  mice per group. Similar data were obtained in a replicate experiment.

vances in this area of parasitology (1, 12, 15, 49). The precise cells first infected and/or acting as antigen-presenting cells (APCs) that signal the adaptive immune system during oral *T. cruzi* infection are not known, but one can envisage at least two possible scenarios as to how the immune system recognizes *T. cruzi* as it progresses from mucosal exposure to a systemic infection. Dissemination could occur early after infection, with trypomastigotes entering the bloodstream soon after invasion or following the first round of replication in cells at the site of initial infection. If this were the case, the phenotype, frequency, and distribution of parasite-specific CD8<sup>+</sup> T cells would be expected to be similar irrespective of the infection route. Alternatively, *T. cruzi* introduced at mucosal surfaces may be restricted to local GI cells for the first cycles of repli-

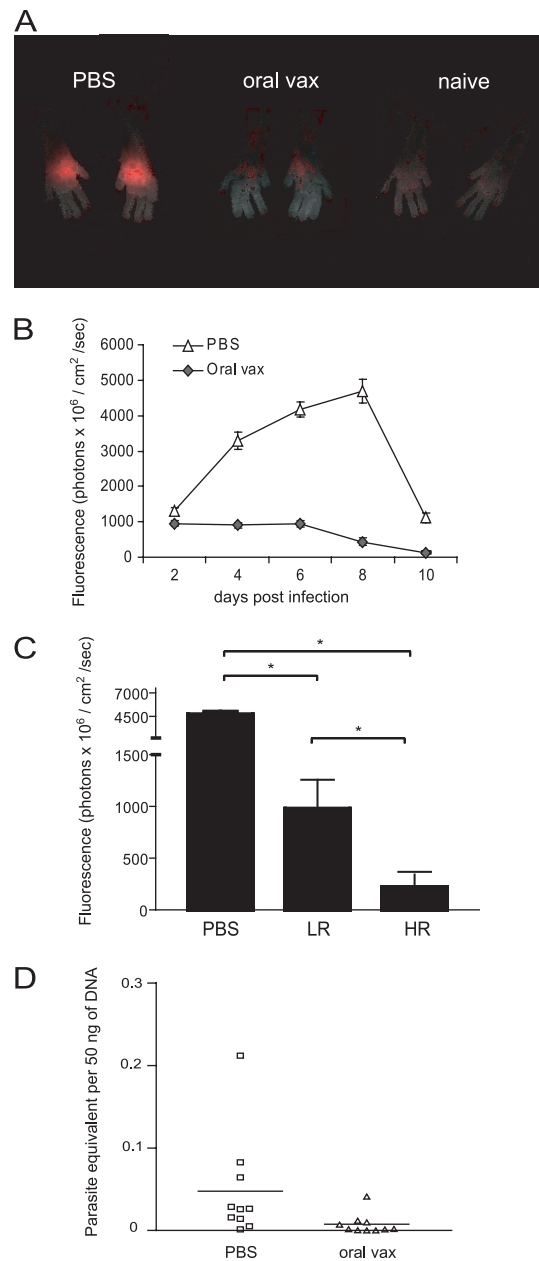
cation and host cell invasion. In this case, Ag presentation would be initially confined to mucosally associated lymphoid tissue, possibly leading to the imprinting of primed T cells for trafficking back to the mucosa. Others have documented local proliferation of parasites following mucosal *T. cruzi* infection, with the early involvement of draining LN, but inflammatory lesions and parasites can be observed at distal sites as early as 14 dpi (21, 26). Our data (Fig. 6) (48) support local parasite amplification following f.p. infection, but parasite DNA is also detectable in dLN within hours of infection (48). These observations suggest that either some parasites migrate out of the infection site prior to replication or that parasites are shuttled to the dLN in lymph or associated with host cells such as DC.

Herein we show that the CD8<sup>+</sup> T cell response to *T. cruzi*



**FIG. 5.** Oral vaccination (vax) with attenuated parasites stimulates *T. cruzi*-specific CD8<sup>+</sup> T cells. (A) Variable degrees of T cell stimulation result from oral vaccination. CD44 expression was assessed 2 weeks after final vaccination on CD8<sup>+</sup> T cells in PBMC. Of 10 vaccinated mice, 3 were designated LR and 7 were designated HR. Flow plots are gated on CD8<sup>+</sup> T cells, and numbers and gates indicate the average percentage of CD44<sup>hi</sup> cells in each group. (B) Oral vaccination generates activated *T. cruzi*-specific CD8<sup>+</sup> T cells. Flow plots are from an individual mouse chosen from HR group to screen for TSKb20<sup>+</sup> responses and CD127 downregulation. All plots are gated on CD8<sup>+</sup> T cells, and numbers indicate the percentage of CD8<sup>+</sup> T cells in a gate or quadrant.

infection develops similarly, regardless of infection route. *T. cruzi*-specific CD8<sup>+</sup> T cells were found in all assayed lymphoid and nonlymphoid (including gut) tissues following either f.p. or p.o. infection. In a different model of oral infection, *T. cruzi* has been shown to replicate in gastric mucosa en route to systemically infecting mice (26). Our initial hypothesis was that replication of parasites in gut-associated tissue such as gastric or intestinal mucosa following oral infection but not f.p. infection may program primed CD8<sup>+</sup> T cells to traffic to gut tissues via expression of  $\alpha 4\beta 7$  (45). However, we were not able to detect differences in the expression of this homing integrin in different routes of infection. The localization of TSKb20-specific cells to GI tissue following f.p. infection was intriguing, given that this compartment often displays stringent “gating” that is permissive only to certain subsets of T cells with a specific homing signature (32). One explanation for the presence of gut-homing *T. cruzi*-specific T cells in f.p.-infected mice is that intestinal tissue is being infected by parasites in the bloodstream. Parasite-induced local inflammation could override the need for  $\alpha 4\beta 7$  expression for T cells to enter gut tissue (3, 53). Alternatively, recent work in a viral infection model suggests that there is a window early in systemic infections during which CD8<sup>+</sup> T cells adopt a gut homing phenotype and populate mucosal sites (36). The protracted course of *T. cruzi* infection



**FIG. 6.** Oral vaccination with attenuated parasites protects mice from WT *T. cruzi* challenge. (A) Parasites are immediately controlled at infection site. Vaccinated mice ( $n = 10$ ) and control mice ( $n = 10$ ) were challenged with  $2.5 \times 10^3$  WT fluorescent *T. cruzi* trypomastigotes injected in superficial subcutaneous tissue of each footpad. Parasite load in f.p. was assessed by quantitating the fluorescent signal with an *in vivo* imaging system. Pictures show left and right feet of individual mice representative of the indicated group at 8 days postchallenge. (B) Vaccine protection is evident at all times through the first 10 days of *T. cruzi* infection. Graph shows the fluorescent signal of all feet in each group at indicated time points (average  $\pm$  SEM,  $P < 0.05$  at all time points,  $n = 20$  feet [10 mice]). (C) Extent of T cell activation is predictive of vaccine efficacy. Parasite load at 8 dpc is graphed for the 4 mice with the highest percentage of CD44<sup>hi</sup> CD8<sup>+</sup> T cells (HR), and the 2 mice with the lowest percentage of CD44<sup>hi</sup> CD8<sup>+</sup> T cells (LR) are graphed along with all mice from PBS group (average  $\pm$  SEM, \* indicates  $P < 0.0001$  by ANOVA). (D) Oral vaccine mice have lower parasite burden in skeletal muscle following WT *T. cruzi* challenge. Parasite load was measured in DNA samples extracted from skeletal muscle of oral vaccine (both HR and LR) and control mice 25 dpc by real-time PCR. Bars show mean of group.



may mean that there is not one particular time during acute infection when a substantial proportion of T cells are expressing  $\alpha 4\beta 7$ . The role of CD103 in the CD8<sup>+</sup> T cell-mediated, adaptive immune response to GI infections is unclear. CD103 appears to be important for retaining T cells in epithelial tissue and dispensable for homing to these sites (19, 36, 59). Though we did not detect a large population of *T. cruzi*-specific CD103<sup>+</sup> CD8<sup>+</sup> T cells in the IEL compartment, it is tempting to speculate that cells with this signature are important for mucosal immunity to *T. cruzi* infection and may contribute to protection against mucosal *T. cruzi* challenge observed by others (23–25, 58). Another possibility is that immunity and protection against mucosal *T. cruzi* challenge are mediated by CD8<sup>+</sup> T cells that transiently survey the gut along with other peripheral tissues. Thus, regardless of route, systemic infection by *T. cruzi* elicits a parasite-specific CD8<sup>+</sup> T cell response in many tissue compartments, including the GI mucosa.

*T. cruzi* also showed tropism for skeletal muscle after oral administration (Fig. 1A) and recently activated (CD69<sup>+</sup>) TSKb20<sup>+</sup> CD8<sup>+</sup> T cells selectively accumulate at that site—an observation also true for f.p.-infected mice (Fig. 2B). Lastly, comparable proportions of *T. cruzi*-specific CD8<sup>+</sup> T cells were activated in lymph nodes draining skeletal muscle or the gut regardless of infection route (Fig. 2D). Overall, these results support a model in which the *T. cruzi*-specific CD8<sup>+</sup> T cell response is initiated by APC that have acquired Ag in sites dispersed throughout the host, consistent with a systemic distribution of parasites beyond the point of parasite entry very shortly after infection.

One reason for undertaking this study was to determine whether we could identify the initial route of infection of hosts by assessing the phenotype of the anti-*T. cruzi* T cells. Although a number of small epidemics of *T. cruzi* infection have clearly been linked to ingestion of contaminated food or drink (14, 28, 67), it is difficult to know the contribution of the oral route of infection when this parasite is acquired in and around houses infested with *T. cruzi*-infected triatomines. Our results suggest that immune parameters will not be particularly useful in determining the mode of transmission in individuals with *T. cruzi* infection, presumably because the infection rapidly becomes systemic irrespective of the route of the initial infection. Another principle reason for studying the immune response following oral infection with *T. cruzi* is that an oral route would be an attractive means of delivery of a vaccine, especially for the protection of reservoir hosts that are integral to the transmission of *T. cruzi* to insect vectors. A number of studies have shown that the interruption of *T. cruzi* transmission to humans could be achieved by decreasing the ability of dogs to transmit *T. cruzi* to insects, a primary factor promoting household infections in many areas where *T. cruzi* is endemic (13, 20, 22). Canine vaccination is a long-standing idea and has a proven role in controlling zoonotic diseases, including rabies and leishmaniasis (40). Reservoirs could conveniently be vaccinated by the oral route, and previous studies support the immunogenic potential of mucosally delivered *T. cruzi* Ag (10, 11, 21, 23–25, 58). It has also been shown that vaccination can generate effective immunity against mucosal *T. cruzi* challenge (23, 24, 58). A key question that had not been addressed is whether a genetically attenuated strain of *T. cruzi* could be used as an oral vaccine.

Here, we show that ECH1<sup>+/-</sup> ECH2<sup>-/-</sup> metacyclic trypomastigotes given by oral gavage provide protective immunity to WT *T. cruzi* challenge. Protection was evident as early as 2 dpc using a new method for tracking the early success of infection by tracking fluorescently tagged parasites (31). Parasite burden at the challenge site was well controlled in vaccinated mice, whereas it increased steadily in controls over the first 8 dpc (Fig. 6B) before declining with the local control of the infection and the dissemination of parasites to other tissues (A. Padilla and R. L. Tarleton, unpublished data). Thus, vaccination reduces the initial “take” of *T. cruzi* infection as well as replication of parasites that successfully invade host cells. The level of CD8<sup>+</sup> T cell activation measured in peripheral blood prior to challenge predicted the degree of protection observed among vaccinated animals. Despite the relative absence of T cell activation in the LR mice, these also exhibit a degree of protection. This result demonstrates that while induction of parasite-specific T cell immunity may be a convenient indicator of vaccine effectiveness, its absence should not absolutely discount a protocol. We chose a heterologous route of challenge in these studies as a more rigorous test of vaccine efficacy. Although there is a discrepancy in the literature as to whether mucosal vaccination is absolutely necessary to induce protection at mucosal sites (4, 31), there is no doubt that mucosal vaccination can stimulate mucosal immunity (27). Additionally, mucosal infection or immunization with *T. cruzi* parasites or Ag leads to IgA production (10, 11, 21, 26, 57) and confers resistance to mucosal *T. cruzi* challenge (23, 24, 58). Thus, given the protection against systemic challenge observed after oral vaccination with *T. cruzi* in these studies, we also predict that an oral route of immunization will effectively protect against mucosal challenge. While strategies such as heterologous prime-boost regimens may maximize immunogenicity for a particular Ag (54), live parasites may best induce a broad-based, protective immune response (63) and be the practical choice to diminish the potential of reservoirs to infect triatomine bugs (22). This work provides support for the further development of attenuated parasite lines for vaccination against *T. cruzi* and defines a system in which to test vaccine candidates and establish correlates of protection.

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