Trypanosoma cruzi Infection Modulates In Vivo Expression of Major Histocompatibility Complex Class II Molecules on Antigen-Presenting Cells and T-Cell Stimulatory Activity of Dendritic Cells in a Strain-Dependent Manner

Catalina D. Alba Soto, Gerardo A. Mirkin, Maria E. Solana, and Stella M. González Cappa*  
Department of Microbiology, Parasitology and Immunology, School of Medicine, University of Buenos Aires, Buenos Aires, Argentina

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A striking feature of Chagas’ disease is the diversity of clinical presentations. Such variability may be due to the heterogeneity among Trypanosoma cruzi isolates or to the host immune response. Employing two strains which differ in their virulence, we investigated the effect of in vivo infection on professional antigen-presenting cells (APC). Acute infection with the virulent RA strain downregulated the expression of major histocompatibility complex (MHC) class II on splenic dendritic cells (DC) and inhibited its induction on peritoneal macrophages and splenic B cells. It also impaired the ability of DC to prime allogeneic T cells and to form homotypic clusters, suggesting a low maturation state of these cells. In contrast, the low-virulence K98 strain maintained the expression of MHC class II on DC or stimulated it on peritoneal macrophages and B cells and preserved DC’s T-cell priming capacity and homotypic clustering. DC from RA-infected mice elicited a lower activation of T. cruzi-specific T-cell proliferation than those from K98-infected mice. APC from RA-infected mice that reached the chronic phase of infection restored MHC class II levels to those found in K98-infected mice and upregulated costimulatory molecules expression, suggesting that the immunosuppression caused by this strain is only transient. Taken together, the results indicate that in vivo infection with T. cruzi modulates APC functionality and that this is accomplished in a strain-dependent manner.

Chagas’ disease, caused by the protozoon Trypanosoma cruzi, affects over 18 million people in Latin America (24). The host-parasite relationship determines the onset of the human pathology, which is extremely diverse, varying from a relatively benign indeterminate form to digestive compromise or a fatal cardiac course (35). Clinical forms might depend on host genetic background and/or intraspecific differences of parasite stocks among different areas: in Argentina, myocardopathy prevails over megacolon, which is the most frequent form in the northeast of Brazil (2, 27).

In order to study the relevance of T. cruzi populations on the host-parasite relationship, we selected two strains with extreme biological differences, RA and CA-I (or its clone K98, with a similar behavior). RA is a highly virulent pantropic/reticulo-tropic strain, while CA-I/K98 is a low-virulence myotropic strain. RA-infected mice develop an early peak of parasitemia during the acute infection, which drops to undetectable values during the chronic phase, whereas CA-I/K98-infected mice slowly increase the parasitemia to reach its highest levels rather late without a sharp drop, remaining measurable for a longer time (13, 14, 26). During the acute stage of the infection, the RA strain is able to induce a transient impairment of both macrophage microbicidal activity and in vitro mitogen-induced lymphoproliferative response. However, CA-I (or the K98 clone) does not affect macrophage microbicidal activity or lymphoproliferative response (5, 6; our unpublished results). At this stage, mice infected with K98 displayed higher prostaglandin E2 serum levels than RA-infected and normal mice. Evidences were provided that this prostaglandin played a role in resistance to infection with the first strain in mice (7). At the chronic phase in RA-infected mice, the nervous system is a privileged target for damage mediated either by CD4 and CD8 T cells or immunoglobulin G antibodies, while for CA-I, only CD4 T cells are able to induce injury to the muscle, which is the target tissue (23, 36).

The activation of T-cell immune responses is mediated by antigen-presenting cells (APC), which deliver antigen-specific and costimulatory signals. Pathogens or their constituents are able to induce APC to deliver these signals to the resting T cells. Some parasites can modulate the antigen presentation and costimulatory activities of APC. Leishmania major induces upregulation of CD40 and CD86 costimulatory molecules (4), but Leishmania amazonensis and Leishmania donovani subvert the APC function of infected macrophages (12, 18). Toxoplasma gondii and Plasmodium falciparum are able to interfere with the processing and presentation of parasite-derived antigen by macrophages and dendritic cells (DC), respectively (22, 37). The persistence of intracellular parasites in immunocompetent hosts suggests that they have evolved multiple strategies to escape or modulate the antigen presentation- and T-cell-mediated parasiticidal activities of the host.

The effect of T. cruzi in vitro infection on macrophages as APC has been documented: Frosch et al. registered a selective upregulation of B7-2 (CD86) costimulatory molecules, although La Flamme et al. proved that a defective T-cell-mac-
 Splenic cell suspensions were simultaneously stained with anti-Mac-3-PE (for peritoneal macrophages) and anti-I-eK-FITC. Peritoneal cells were simultaneously stained with anti-CD11c-PE (for DC) and anti-I-eK-FITC. Spleen cell suspensions were simultaneously stained with anti-CD11c-PE (for B cells) and anti-I-eK-FITC. CD11c+, Mac-3+, and CD19+ events were analyzed for I-eK expression. The mean fluorescence intensity of cells with specific conjugated monoclonal antibody was <10. Results of one representative experiment out of five are shown.

rophere adhesion impaired antigen presentation (11, 20). The parasite population employed in each assay could be responsible for these discordant findings. Van Overveldt et al. demonstrated that in vitro T. cruzi infection prevents monocyte-derived human DC from optimal maturation and expression of HLA-DR molecules (38). However, no information is available about the in vivo interaction of APC with T. cruzi in vivo.

In the present study, we investigated the influence of in vivo infection with T. cruzi strains which differ in virulence on professional APC expression of major histocompatibility complex (MHC) class II and costimulatory molecules and on the T-cell stimulatory capacity of spleenic DC.

**MATERIALS AND METHODS**

**Mice.** Male C3H/HeN and C57BL/6 mice were supplied by the animal facilities of the Department of Microbiology, School of Medicine, University of Buenos Aires.

**Parasites, infection, and T. cruzi lysate.** Bloodstream forms (trypomastigotes) of the RA strain or the K98 clone (derived from the CA-1 strain) of T. cruzi were maintained by serial passages in mice. At 8 weeks of age, C3H/HeN mice were infected with 100 trypomastigotes of either T. cruzi population. Except when indicated, trypomastigotes were inoculated into the hind footpad. By this route, at least 40% of animals infected with the virulent RA strain reached the chronic phase of infection. Uninfected mice of the same age, body weight, and sex were used as controls. For studies at the acute phase of infection, mice were killed when circulating trypanosomes were detected by the microhematocrit technique (10) (15 days for RA-infected and 20 days for K98-infected mice) and 4 months postinfection for those at the chronic phase.

**Trypanosomes** of the RA and K98 T. cruzi strains were purified from whole blood by density gradient centrifugation as described before (25). Trypomastigotes were subjected to five freeze-thawing cycles, resuspended in phosphate-buffered saline and sonicated (10 cycles of 30 s each at 40 Hz on ice). Suspensions were centrifuged, and the supernatant was filtered and stored at −20°C until use. Protein concentration was determined by the method of Lowry et al. with bovine serum albumin (Sigma, St. Louis, Mo.) as the standard.

**Cells.** To prepare dendritic cell (DC)-enriched suspensions, spleens were homogenized at 4°C in a tissue grinder with RPMI 1640 supplemented with 10% fetal bovine serum (FBS), 2-mercaptoethanol, and antibiotics (penicillin at 100 U/ml and streptomycin at 100 μg/ml (Sigma). Homogenates were subjected to digestion with collagenase (1 mg/ml), DNase I (1 mg/ml), and hyaluronidase (0.5 mg/ml) (Sigma) for 30 min at 37°C. Undigested stromal fragments were removed, and the cells, resuspended in 1.080-g/ml density bovine serum albumin solution, were centrifuged at 4,500 × g for 25 min. The low-density fraction was recovered and incubated at 37°C for 2 h in RPMI 1640–10% FBS. Nonadherent cells were removed, and the remaining ones were reincubated in RPMI 1640–10% FBS for 20 h, the time required for DC to detach from plastic.

To purify splenic DC, CD11c+ cells were isolated by magnetic sorting with biotinylated anti-CD11c (HL3) monoclonal antibodies bound to streptavidin-coated magnetic beads and MiniMACS columns (Miltenyi Biotech GmbH, Bergisch Gladbach, Germany) following the manufacturer’s protocol. Cell populations obtained in this way consisted of approximately 90% CD11c+ cells as determined by flow cytometry analysis.

Peritoneal cells were recovered by intraperitoneal injection of 5 ml of cold RPMI 1640 supplemented with heparin (5 U/ml). After gentle abdomen massage, the fluid was aspirated, and cells were washed twice with RPMI–10% FBS and kept at 4°C until use.

**Flow cytometry.** Cells were washed twice in ice-cold phosphate-buffered saline supplemented with 1% bovine serum albumin and 0.1% Na2EDTA and resuspended at a final cell concentration of 2 × 10⁶/ml. Two percent normal mouse serum was added to avoid unspecific binding through Fc receptors. Then, cells were incubated for 30 min at 4°C with previously optimized amounts of one or more of the following conjugated murine monoclonal antibodies: anti-CD11c-phycocerythrin (PE), anti-CD19-PE, anti-Mac-3-PE, anti-I-eK-PE (MHC II)-fluorescein isothiocyanate (FITC), anti-CD4-FITC, anti-CD8-IFITC, and the biotinylated monoclonal antibodies anti-CD24, anti-CD40, anti-CD80, and anti-CD86. All monoclonal antibodies were purchased from PharMingen, San Diego, Calif. Streptavidin-Cy-chrome (PharMingen) was used as a second-step reagent. As controls, cells were stained with the corresponding isotype-matched monoclonal antibody.

**Cells were fixed with 1% paraformaldehyde before being acquired with a flow cytometer (Ortho Cytomront Absolute; Johnson & Johnson). Data were analyzed with the WINMDI software (Joseph Trotter, Scripps Research Institute). Flow cytometry results were expressed as the median fluorescence intensity (median fluorescence intensity of cells with specific monoclonal antibody – median fluorescence intensity of isotype-matched monoclonal antibody):**

**Trypanosoma inhabits human and rodent macrophages (21, 22) and expresses a variety of surface molecules (23). Infection of these cells results in a...**

**Flow cytometry was performed as described above. Data were analyzed with the WINMDI software (Joseph Trotter, Scripps Research Institute). Flow cytometry results were expressed as the median fluorescence intensity (median fluorescence intensity of cells with specific monoclonal antibody – median fluorescence intensity of isotype-matched monoclonal antibody):**

**TABLE 1. Surface expression of I-eK (MHC class II) molecules on professional APC from T. cruzi-infected mice**

<table>
<thead>
<tr>
<th>Strain, inoculation route, and phase</th>
<th>CD11c+ cells</th>
<th>Mac-3+ cells</th>
<th>CD19+ cells</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean MFI ± SD</td>
<td>Mean % I-eK+ ± SD</td>
<td>Mean MFI ± SD</td>
</tr>
<tr>
<td>Control</td>
<td>1,242 ± 73</td>
<td>78.5 ± 3.9</td>
<td>325 ± 62</td>
</tr>
<tr>
<td>K98, FPI, acute</td>
<td>1,026 ± 160</td>
<td>83.3 ± 3.5</td>
<td>1,344 ± 140**</td>
</tr>
<tr>
<td>RA, FPI, acute</td>
<td>489 ± 29**</td>
<td>11.3 ± 7.3***</td>
<td>349 ± 176</td>
</tr>
<tr>
<td>K98, PI, acute</td>
<td>1,111 ± 161</td>
<td>73.0 ± 6.0</td>
<td>1,022 ± 249*</td>
</tr>
<tr>
<td>RA, PI, acute</td>
<td>532 ± 238*</td>
<td>31.3 ± 9.3***</td>
<td>441 ± 38</td>
</tr>
<tr>
<td>K98, FPI, chronic</td>
<td>1,309 ± 86</td>
<td>95.5 ± 1.5</td>
<td>1,134 ± 240**</td>
</tr>
<tr>
<td>RA, FPI, chronic</td>
<td>1,900 ± 136**</td>
<td>83.0 ± 3.0</td>
<td>1,796 ± 95**</td>
</tr>
</tbody>
</table>

* Mice were infected with RA or K98 strain by the foot pad (FPI) or peritoneal (PI) route of inoculation. Cells were obtained at the acute or the chronic phase of infection. CD11c+ (for DC), Mac-3+ (for peritoneal macrophages) and CD19+ (for B cells) events were analyzed for I-eK-FITC expression. Results are expressed as the mean fluorescence intensity (MFI) of I-eK expression or mean percentage of I-eK+ cells. *, P < 0.05; **, P < 0.01; and ***, P < 0.001 compared with the control.
Fluorescence intensity of cells with irrelevant isotype-matched monodonal antibody or as the percentage of cells which were positive for the surface marker evaluated.

**Proliferation assays.** For the allogeneic mixed-lymphocyte reaction, irradiated (20 Gy) splenic DC from infected and uninfected mice were cocultured with untreated CD4 or CD8 T cells from the lymph nodes of C57BL/6 uninfected mice at 37°C in RPMI 1640–10% FBS medium in 96-well microplates in triplicate at a final volume of 200 μl. For evaluation of T. cruzi-specific T-cell responses, irradiated splenic DC from infected and uninfected mice were cocultured with untreated CD4 T cells from the lymph nodes of chronically infected mice and controls in the presence of T. cruzi lysate (50 μg/ml) at 37°C in RPMI 1640–10% FBS medium in 96-well microplates in triplicate at a final volume of 200 μl. Cultures were pulsed with 0.5 μCi of [3H]thymidine (ICN, Costa Mesa, Calif.) per well during the last 24 h of culture. Cells were harvested and analyzed in a Rack Beta liquid scintillation counter (Pharmacia). Results are expressed as the mean counts per minute (cpm) of the culture ± standard deviation.

Statistical analysis. One-way analysis of variance and Bonferroni’s multiple comparison test were used for the analysis of statistical significance.

**RESULTS**

**Effect of infection with T. cruzi strains RA and K98 on surface expression of MHC class II and costimulatory molecules of professional APC.** DC located in lymphoid organs constitutively express large amounts of MHC class II molecules at their surface; macrophages and B cells upregulate MHC class II expression in response to infection (1, 28). Flow cytometry analysis showed that constitutive expression of MHC class II in DC was downregulated during the acute phase in mice infected with strain RA while in peritoneal macrophages and B cells, an inhibition of the infection-enhanced MHC class II expression was registered in the same group of animals (Fig. 1). This phenomenon was observed not only as a decrease in the percentage of positive cells but also as a reduction in the number of surface molecules per cell (Table 1). In contrast, MHC class II expression was not altered on DC by infection with strain K98 (Fig. 1A) and was adequately upregulated on peritoneal macrophages (Fig. 1B) and B cells (Fig. 1C). This strain-dependent effect of T. cruzi infection on professional APC was independent of the route of inoculation, since similar results were obtained when the intraperitoneal or the footpad route was used to infect animals (Table 1). It is noteworthy that APC from RA-infected mice who reached the chronic stage of infection attained or even surpassed the levels of MHC class II expression reached by APC from K98-infected mice (Table 1).

As shown in Table 2, surface expression of CD86 on splenic DC was downregulated by acute infection independently of the T. cruzi population employed. Interestingly, at the chronic phase, peritoneal macrophages and B cells of mice surviving the acute phase of infection with RA showed significantly higher levels of CD86 expression than those from K98-infected mice. This was also the case for DC from chronically RA-infected mice, whose CD80 expression was higher than in K98-infected animals. T. cruzi infection had no effect on the basal expression of CD40 ad CD24 molecules (data not shown).

**Effect of acute infection with T. cruzi RA and K98 strains on distribution of APC in spleens.** APC settle in secondary lymphoid organs, where they encounter T cells for antigen presentation (1). As shown in Table 3, flow cytometry analysis of the relative number of professional APC located in the spleen revealed no significant differences in the percentages of CD11c+ (DC) cells among the three groups studied. However, an increase in the percentage of Mac-3+ cells in the spleens of RA-infected mice and a reduction in the relative number of CD19+ cells were found in the same group of animals. This last finding in spleens from RA-infected mice is consistent with the high rate of spontaneous apoptosis of splenic B cells by infection with the virulent Tulahuen strain of T. cruzi reported by Zuñiga et al. (40).

**Acute infection with the RA T. cruzi strain drastically diminishes DC stimulation of CD4 T and CD8 T-cell alloresponse.** The ability of APC to stimulate an alloresponse strongly depends on the first signal provided by MHC class II molecules. Here we employed DC, the most potent inducers of T-cell immune responses in vitro, to test this capacity on professional APC from T. cruzi-infected mice (3). We performed a mixed-lymphocyte reaction with irradiated DC from acutely RA- or K98-infected and control C3H/HeNk mice as stimulators and purified lymph node CD4 T cells from C57BL/6 mice as responders. Allogeneic CD4 T cells cocultured with DC from control and K98-infected mice proliferated readily, whereas those cocultured with DC from RA-infected mice were unresponsive (Fig. 2). DC are equally important in priming naive CD8 T cells. They can stimulate the proliferation of naive CD8 T cells.

**TABLE 2. Surface expression of costimulatory molecules on professional APC from T. cruzi-infected mice**

<table>
<thead>
<tr>
<th>Strain and infection phase</th>
<th>CD11c+ cells</th>
<th>Mac-3+ cells</th>
<th>CD19+ cells</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>CD80</td>
<td>CD86</td>
<td>CD80</td>
</tr>
<tr>
<td></td>
<td>1,504 ± 174</td>
<td>1,453 ± 124</td>
<td>1,640 ± 76</td>
</tr>
<tr>
<td>K98, acute</td>
<td>1,196 ± 99</td>
<td>928 ± 68*</td>
<td>1,677 ± 48</td>
</tr>
<tr>
<td>RA, acute</td>
<td>1,457 ± 172</td>
<td>1,063 ± 24*</td>
<td>1,458 ± 146</td>
</tr>
<tr>
<td>K98, chronic</td>
<td>1,298 ± 141</td>
<td>1,179 ± 19</td>
<td>1,795 ± 304</td>
</tr>
<tr>
<td>RA, chronic</td>
<td>1,846 ± 44**</td>
<td>1,299 ± 225</td>
<td>1,930 ± 149</td>
</tr>
</tbody>
</table>

* Mice were infected with RA or K98 T. cruzi strains by the footpad (FPI) route of inoculation. CD11c+, Mac-3+, and CD19+ events were analyzed for CD80 and CD86 expression. *, P < 0.05 compared with controls; **, P < 0.05 comparing K98 chronic and RA chronic.

**TABLE 3. Relative number of APC in spleens from mice during the acute phase of infection with T. cruzi**

<table>
<thead>
<tr>
<th>Infection</th>
<th>Mean % of cells positive (± SD) for surface marker:</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>CD11c+</td>
</tr>
<tr>
<td>Control</td>
<td>3.4 ± 0.4</td>
</tr>
<tr>
<td>RA</td>
<td>6.4 ± 1.0</td>
</tr>
<tr>
<td>K98</td>
<td>4.1 ± 1.2</td>
</tr>
</tbody>
</table>

* Mice were infected with the RA or K98 T. cruzi strain by footpad inoculation. Splenic single-cell suspensions were stained for PE-conjugated monoclonal antibodies against the cell surface markers. Results are expressed as mean percent positive cells. *, P < 0.05; **, P < 0.01 compared with the control.
allogeneic CD8 T cells in the absence of CD4 T-cell cooperation (16). In another set of experiments with CD8 T cells instead of CD4 T cells, a mixed-lymphocyte reaction was performed to test this trait. DC from RA-infected mice were less efficient at priming allogeneic CD8 T cells for the proliferation response (Fig. 2).

In addition to modulating the allostimulatory capacity of DC, pathogens and pathogen-derived factors themselves can drive DC to produce or induce the release of cytokines promoting different types of immune responses (30). To establish whether DC from RA-infected mice were producing or inducing the production of factors which limited T-cell proliferation (e.g., transforming growth factor beta or interleukin-10), we tested whether addition of DC from uninfected mice (DC control) to the mixed-lymphocyte reaction of DC from RA-infected mice plus CD4 T cells from C57BL/6 mice for the last 3 days of culture would reconstitute the T-cell alloreponse. However, as shown in Fig. 3, culture alloreponse was not restored on the sixth day of culture to the levels found with control DC. Furthermore, addition of supernatant from the mixed-lymphocyte reaction of DC from RA-infected mice with C57BL/6 (C57) CD4 T cells (supDCRA). Background proliferation of DC alone was ≤400 cpm and of CD4 T cells alone was ≤500 cpm. Cell proliferation was measured after a 24-h pulse with [3H]thymidine. Values are the means of triplicate cultures ± standard deviation. Results for one of three independent experiments are shown.

Effect of in vivo infection with T. cruzi on ability of DC to activate T. cruzi-specific T-cell proliferative responses. We investigated ex vivo the antigen-presenting activity of DC isolated from the spleens of infected and control mice and monitored their capacity to stimulate CD4 T cells isolated from infected and control mice. As shown in Fig. 4, DC recovered from the spleens of K98-infected mice mediated the highest proliferation of CD4 T cells from infected and control animals. However, stimulation was more efficient with CD4 T cells obtained from mice chronically infected with the homologous T. cruzi strain. In contrast, DC recovered from RA-infected mice induced a weaker proliferation of CD4 T cells than DC from K98-infected mice.

Splenitic DC from acutely RA-infected mice show features of low state of maturation. Following antigen capture, immature DC migrate to lymphoid organs, where, after maturation, they display peptide-MHC complexes (3, 15). Homotypic clustering of DC is a close correlate of their state of maturation, as it reflects morphological and physiological changes, including rearrangements of adhesion molecules at the cell surface (8, 34).
Here, isolated splenic DC obtained from acutely T. cruzi of infection with RA T. cruzi on the effect on professional APC of infection in vivo.

Alterations of APC infected in vitro with type and state of APC activating T cells are of great importance are still being investigated. Nevertheless, it is clear that the from RA-infected mice formed fewer and smaller DC-DC clusters. Isolated splenic DC from mice infected with the K98 strain (A) or RA strain (B) at the acute phase of infection are shown. Cells were cultured in RPMI-10% FBS for 12 h at 37°C. Bar, 20 μm.

**FIG. 5.** Cultured splenic DC obtained from mice at the acute phase of infection with RA T. cruzi strain are unable to form homotypic cell clusters. Isolated splenic DC from mice infected with the K98 strain (A) or RA strain (B) at the acute phase of infection are shown. Cells were cultured in RPMI-10% FBS for 12 h at 37°C. Bar, 20 μm.

Here, isolated splenic DC obtained from acutely T. cruzi-infected mice were cultured for 12 h to test this trait. DC isolated from RA-infected mice formed fewer and smaller DC-DC clusters (Fig. 5B) than DC from K98-infected mice (Fig. 5A) or controls (not shown).

**DISCUSSION**

The mechanisms of immune response against T. cruzi infection are still being investigated. Nevertheless, it is clear that the type and state of APC activating T cells are of great importance. Alterations of APC infected in vitro with T. cruzi have been communicated (11, 20, 29, 38), but there are no reports on the effect on professional APC of infection in vivo.

In the present study, we show that murine infection with T. cruzi modulates the expression of MHC class II molecules on professional APC at their plasma membrane in a strain-dependent manner. Indeed, infection with the low-virulence K98 strain preserved the expression of this molecule on DC or stimulated it on peritoneal macrophages and B cells, while the virulent RA strain reduced or inhibited its expression at the surface of professional APC. In the in vivo infection, we found occasional DC and macrophages harboring amastigotes by immunofluorescence staining (less than 5%) without significant differences between strains in the parasite load of APC (data not shown). These results show that factors other than infection of APC themselves are involved in downregulating MHC class II expression during the acute phase of infection with the virulent RA strain. Our results obtained in vivo seem to corroborate what Van Overtvelt et al. demonstrated in vitro: close contact between DC and T. cruzi is not a prerequisite for alteration of the optimal maturation of human DC and expression of HLA-DR molecules (38).

The RA strain almost abrogates the ability of DC to prime allogeneic CD4 T cells, suggesting that the low MHC II expression is responsible for this alteration. Furthermore, DC from RA-infected animals lacked the ability to stimulate alloreactive CD8 T cells, indicating a poor functional capability of these cells. Interestingly, in the mixed-lymphocyte reaction, alloresponse was not restored by addition of DC from uninfected mice to cultures containing DC from RA-infected animals, and supernatants of DC from RA-infected mice inhibited T-cell allostimulation. This finding suggests that infection with the virulent RA strain could be favoring the induction or production of suppressive/regulatory cytokines by DC, thus driving T cells to anergy or to a regulatory phenotype. Even though MHC class II-dependent presentation by DC was not completely lost after infection, DC from K98-infected mice induced a stronger T. cruzi-specific T-cell response than those from RA-infected mice. The fact that DC from RA-infected mice can still elicit a noticeable T-cell response even with low MHC class II expression is not unexpected because it has been proved that only small amounts of MHC class II are necessary to mount a functional cellular response in vivo (39).

DC homotypic interactions stimulate the accessory function of DC by mutual delivery of maturation signals and transfer of antigen between cells (8). The reduced DC-DC clustering of DC from RA-infected mice, together with the low MHC class II expression, might point to an interference with maturation process induced by acute infection with this virulent strain. It has been reported that antigen presentation by immature DC might induce the differentiation of naive T cells towards a suppressor regulatory phenotype capable of interfering with DC-mediated Th1 differentiation (33). Our observations add the virulent RA strain of T. cruzi to a growing list of stimuli, including measles virus and herpesvirus and *Plasmodium falciparum*-infected erythrocytes, which appear to inhibit terminal DC differentiation at various stages (17, 34). In contrast, infection with the low-virulence K98 strain preserved DC maturation and allogeneic or T. cruzi-specific T-cell response, allowing the host to control the parasitemia and to survive the acute infection.

The CD80/CD86-CD28 costimulatory pathway participates in the pathogenesis of several infectious diseases (31). As CD86 is the most critical for amplification of T-cell responses (9, 21), its downmodulation on splenic DC from infected mice might contribute to the depression of immune responses observed in acute infection with T. cruzi, a phenomenon of great importance in the pathogenesis of Chagas’ disease (19).

It is of interest that APC from mice infected with the virulent RA strain who survived the acute phase of infection recovered or even surpassed MHC class II and costimulatory molecule expression levels to those of K98-infected mice. This finding might be correlated with the kinetics of T- and B-cell proliferation because unresponsiveness to polyclonal activators is seen only at the acute phase of infection with the virulent RA strain. Our data suggest that alteration of APC function by infection with the virulent RA strain is a transient phenomenon during acute infection.

Recent studies reveal that induction of an immune response to pathogens is determined by several variables, including DC function itself, pathogen-derived signals, the local microenvironment, and cytokines released by neighboring T cells and other cells (30, 32). Our results could reflect strain-dependent differences in these microenvironments. Functional modulation of APC by the virulent RA strain may establish an environment that contributes to the great susceptibility shown during acute infection with this strain. An analysis of the immune response focused on the cytokine profiles is currently in progress in our laboratory.
In vitro models are useful to analyze the interactions between APC and *T. cruzi*. However, the actual relevance of these interactions needs to be demonstrated in vivo, something that has not been done so far. Our results put in evidence for the first time that *T. cruzi* infection can modulate APC function in vivo. Furthermore, we found that the degree of this alteration correlates with the virulence of the *T. cruzi* population infecting the host.

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