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Levels of expression of costimulatory molecules have been proposed to influence the outcome of antigen-specific T cell priming. We found that *Leishmania major* selectively modulated the expression of costimulatory molecules on various populations of epidermal cells. B7.2 expression was down-regulated on Thy1.2⁺ epidermal cells (keratinocytes) from disease-resistant C3H mice, but not from disease-susceptible BALB/c mice. In addition, epidermal cells from BALB/c mice showed a down-regulation of B7.1 expression on NLDC 145⁺ Langerhans cells. *In vivo* T cell priming experiments, using syngeneic epidermal cells as antigen-presenting cells (APC), showed that the production of IFN- γ was inhibited when either B7.1 or B7.2 signaling pathways were blocked. Blockade of B7.2, but not B7.1, significantly inhibited the ability of epidermal cells to induce IL-4 production from CD4⁺ T cells. In addition, C3H CD4⁺ T cells, which were unable to secrete detectable levels of IL-4 in cultures with syngeneic APC, were now able to secrete IL-4 following presentation of *L. major* antigens by congenic BALB/K epidermal cells. Conversely, C3H epidermal cells supported the priming of BALB/K CD4⁺ T cells for IL-4 production *in vitro*. Thus, the differential expression of B7 molecules on epidermal cells may not represent the sole factor governing the polarization of *L. major*-specific CD4⁺ T cells *in vivo*.

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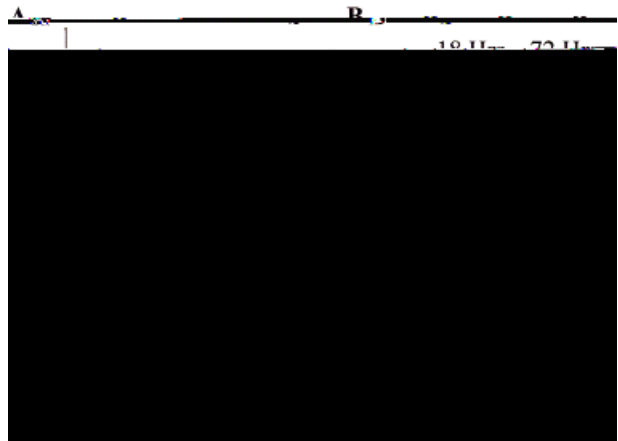
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The protozoan parasite *Leishmania major* induces cutaneous leishmaniasis. It exists as a flagellated promastigote in its insect vector, the sand fly. The vertebrate host becomes infected with *L. major* when the sand fly probes into the skin for a blood meal and injects parasites. Promastigotes are taken up by phagocytic cells, and within these cells

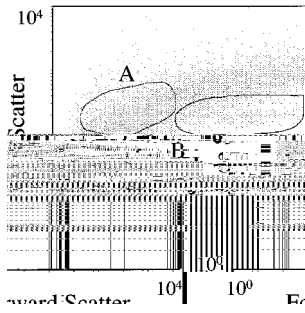
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Abb a LC: Langerhans cells L : *Leishmania major* MFI: Mean fluorescence intensity

Because the skin represents the primary target organ of *L. major*, epidermal APC, such as Langerhans cells (LC), are likely to play a major role in the initiation of an immune response against *L. major*. In fact, previous studies have shown that epidermal LC, but not keratinocytes, present *L. major* antigens to T cells [28–30]. Fur-

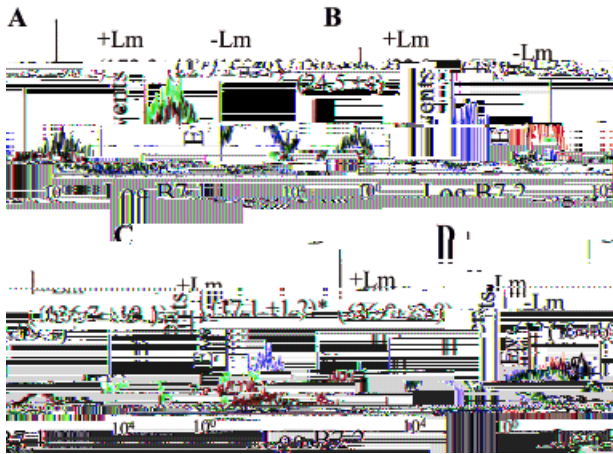


Fi Up-regulation of B7 molecules on epidermal cells cultured in the absence of *L. major*. Epidermal cells from (A, C) BALB/c and (B, D) C3H mice were analyzed for B7.1 using PE-conjugated anti-B7.1 and B7.2 expression using FITC-conjugated anti-B7.2 at 18 h (blue lines) and 72 h (black lines). Red lines represent isotype control Ab. For clarity, only isotype control Ab using cells cultured without *L. major* are shown. Similar results were obtained when epidermal cells cultured with *L. major* were stained with the same isotype control Ab. The data are representative of three experiments.

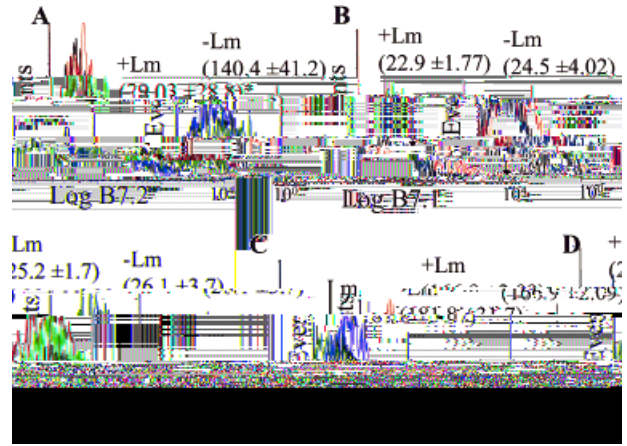


Fi Light scatter gates of NLDC 145 and Thy1.2⁺ BALB/c epidermal cells. Epidermal cells were double-stained with NLDC-145 and anti-Thy1.2 mAb. The gates shown enclose NLDC 145⁺ (A) and Thy1.2⁺ (B) epidermal cells. No NLDC 145⁺/Thy1.2 double-positive cells were identified. The light scatter gates shown here were used in subsequent experiments to enclose populations containing NLDC 145⁺ and Thy1.2⁺ cells.

(light scatter gates of the different epidermal cell populations are shown in Fig. 2). Both B7.1 and B7.2 were equally up-regulated on NLDC 145⁺ epidermal cells (LC) from C3H mice in the presence or absence of *L. major* (Fig. 3A, B; *p* values in each figure were greater than 0.2). However, *L. major* induced a selective down-regulation of B7.1 expression on BALB/c NLDC 145⁺ cells [Fig. 4A; mean fluorescence intensity (MFI) of 140±41.2 in the

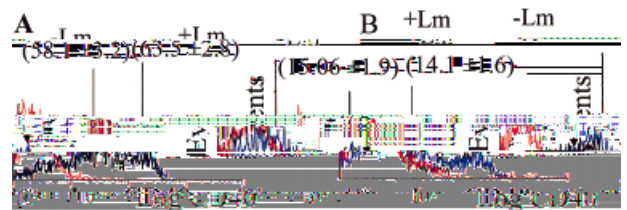


Fi *L. major* selectively down-regulates B7.2 expression on Thy1.2⁺ epidermal cells from C3H mice. Epidermal cells were isolated and cultured with (black lines) or without (blue lines) *L. major* (Lm) promastigotes and analyzed by flow cytometry 3 days later by gating on (A, B) NLDC 145⁺ or (C, D) Thy1.2⁺ C3H epidermal cells predetermined by light scatter. Red lines represent isotype control Ab. Numbers in parentheses indicate MFI ± SD from three experiments. The data are representative of three experiments. **p*=0.03.



Fi *L. major* selectively down-regulates B7-1 expression on BALB/c NLDC 145⁺ LC. Epidermal cells were isolated and cultured with (black lines) or without (blue lines) *L. major* (Lm) promastigotes and analyzed by flow cytometry 3 days later by gating on (A, B) NLDC 145⁺ or (C, D) Thy1.2⁺ BALB/c epidermal cells predetermined by light scatter. Red lines represent isotype control Ab. Numbers in parentheses indicate MFI ± SD from three experiments. The data are representative of three experiments. **p*=0.05.

absence of *L. major* compared to 179±28.8 in the presence of *L. major*, *p*=0.05]. In contrast, B7.2 expression on BALB/c NLDC 145⁺ was not affected in cultures containing *L. major* (Fig. 4B). Our results also showed that *L. major* differentially regulates the expression of B7.1 and B7.2 on Thy1.2⁺ epidermal cells (e.g. keratinocytes and dendritic epidermal T cells). B7.2 expression on Thy1.2⁺ C3H epidermal cells was down-regulated in the presence of *L. major* (Fig. 3D; MFI of 36.9±2.87 in the absence of *L. major* compared to a MFI of 17.1±1.2 in the presence of *L. major*, *p*=0.03). *L. major* did not alter



Fi Equal up-regulation of CD40 expression on NLDC 145⁺ epidermal cells derived from BALB/c and C3H mice in the presence or absence of *L. major*. Epidermal cells derived from (A) C3H and (B) BALB/c mice were cultured in the presence (black lines) or absence (blue lines) of *L. major* (Lm) and CD40 expression was analyzed 3 days later. Light scatter gates were predetermined to enclose NLDC 145⁺ cells. Red lines represent isotype control Ab. Numbers in parentheses indicate intensity MFI ± SD from three experiments. The data are representative of three experiments.

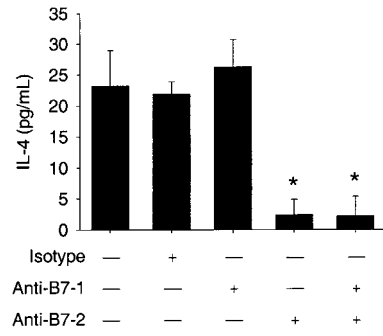
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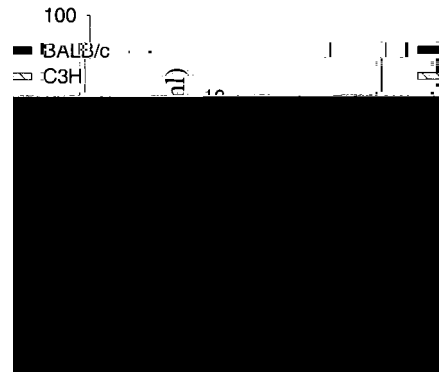
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Fi B7.2, but not B7.1, on BALB/c epidermal cells costimulates for IL-4 production by *L. major*-specific CD4⁺ T cells



Tab 1. *In vivo* production of IL-4 using a congenic C3H-BALB/K mouse model^{a)}

APC-T cells	IL-4 (pg/ml)				
	No Ab	Isotype	anti-B7.1	anti-B7.2	anti-B7.1 + anti-B7.2
C3H-C3H	0	0	0	0	0
C3H-BALB/K	42.6 ± 5.1	35.2 ± 3.4	45.8 ± 6.2	10.4 ± 2.2*	3.1 ± 0.75*
BALB/K-BALB/K	52.1 ± 3.8	55.7 ± 4.3	59 ± 6.8	8.6 ± 3.5*	2.4 ± 0.65*
BALB/K-C3H	69 ± 4.3	55 ± 7.4	58.8 ± 7.1	6.7 ± 1.2*	2.9 ± 0.32*

a) The cultures were set up as described in Fig. 6 and Sect. 4. The data represent means ± SD from three experiments. * $p < 0.05$.

Tab 2. *In vivo* production of IFN- γ using a congenic C3H-BALB/K mouse model^{a)}

APC-T cells	IFN- γ (ng/ml)				
	No Ab	Isotype	anti-B7.1	anti-B7.2	anti-B7.1 + anti-B7.2
C3H-C3H	5.8 ± 1.3	4.1 ± 1.8	2.8 ± 0.5*	1.9 ± 0.78*	0.29 ± 0.1*
C3H-BALB/K	4.6 ± 2.3	5.3 ± 1.7	3.2 ± 1.1*	2.5 ± 1.05*	0.35 ± 0.2*
BALB/K-BALB/K	6.9 ± 1.7	6.3 ± 2.2	3.4 ± 1.4*	2.9 ± 1.5*	0.28 ± 0.17*
BALB/K-C3H	5.7 ± 2.3	6.1 ± 1.8	2.8 ± 0.8*	3.3 ± 1.9*	0.4 ± 0.18*

a) The cultures were set up as described in Fig. 7 and Sect. 4. The data represent means ± SD from three experiments. * $p < 0.05$.

2.4 *In vivo* production of IL-4 and IFN- γ in a congenic BALB/K-C3H mouse model

To determine whether factors other than B7 expression on epidermal cells influence cytokine production by CD4⁺ T cells, we used a congenic BALB/K (disease-susceptible)-C3H (disease-resistant) model to analyze IL-4 and IFN- γ production *in vivo*. As shown in Table 1, epidermal cells from C3H mice were able to prime BALB/K CD4⁺ T cells for IL-4 production. Conversely, BALB/K epidermal cells supported the production of IL-4 by CD4⁺ T cells from C3H cells. In all cases, anti-B7.2, but not anti-B7.1 mAb, significantly inhibited the generation of IL-4 *in vivo*. These results suggest that costimulation for IL-4 production by CD4⁺ T cells *in vivo* may be regulated by more than costimulatory molecules on epidermal cells since BALB/K T cells produced IL-4 when activated by either BALB/K T cells or C3H epidermal cells.

The production of IFN- γ was also assessed in the congenic model described above. Both C3H and BALB/K epidermal cells were able to prime CD4⁺ T cells from BALB/K and C3H mice, respectively (Table 2). Consistent with the data reported in the syngeneic model (Fig. 7), the generation of IFN- γ was dependent on both B7.1 and

B7.2 molecules, because addition of anti-B7.1 or anti-B7.2 to the cultures significantly inhibited the secretion of IFN- γ (Table 2).

3 Discussion

The T cell cytokine profile elicited in the host represents a crucial factor in determining disease outcome in mice infected with *L. major* [2–5]. During primary T cell responses, the pattern of cytokines elicited by Ag-specific T cells may be regulated, at least in part, by the strength and affinity of the interaction between accessory molecules and their coreceptors on T cells [19, 20, 22]. Therefore, the levels of expression of accessory molecules on APC able to initiate a primary immune response could represent a crucial factor in determining the outcome of T cell priming. Because the skin repre-

Table 3). On BALB/c epidermal cells, B7.2 expression was equally up-regulated on Thy-1.2⁺ and NLDC-145⁺ cells in the presence or absence of *L. major*, whereas B7.1 expression was down-regulated on NLDC 145⁺ cells (Fig. 4A). Furthermore, the B7.2 signaling pathway on BALB/c epidermal cells appeared to be involved in IL-4 production by *L. major*-specific CD4⁺ T cells, because addition of neutralizing anti-B7.2, but not anti-B7.1, mAb significantly reduced the levels of IL-4 produced (Fig. 6). These results suggest that costimulation via B7.2 on LC could promote the production of IL-4 in susceptible mice infected with *L. major*. This hypothesis is supported by previous findings showing that treatment of BALB/c mice with neutralizing anti-B7.2 mAb dramatically reduced the levels of IL-4 produced in the lymph nodes draining leishmanial lesions, and enhanced resistance to *L. major* infection [24]. Likewise, Corry et al. [23] reported that treatment of BALB/c mice with CTLA4Ig within the first week of infection completely abrogated progressive disease, suggesting that the priming of Th2

strated in studies showing that CD40-deficient mice on a resistant background were unable to control infection with *L. major* [25, 26]. Thus, we compared levels of CD40 expression on BALB/c and C3H epidermal cells. Both strains of mice showed up-regulation of CD40 expression on NLDC 145⁺ cells to equal maximum levels in the presence or absence of *L. major*. However, a broader range of CD40 expression was observed on C3H cells compared to BALB/c cells (Fig. 5). It is unclear how this difference in the population of CD40⁺ epidermal cells impacts the outcome of T cell priming, but further study to examine this issue is warranted.

The selective down-regulation of B7.2 on Thy-1.2⁺ positive C3H epidermal cells (Fig. 3D) argues for potential important roles played by Thy-1.2⁺ epidermal cells, such as keratinocytes, in cutaneous leishmaniasis. Since

4.4 Infection of epidermal cells

Freshly isolated LC-enriched epidermal cells were resuspended in DMEM (Gibco BRL) supplemented with 10% heat-inactivated FBS (Hyclone, Logan, UT), 0.1 mM nonessential amino acids (Gibco), 0.05 mM 2-mercaptoethanol, 10 mM HEPES (Gibco), 1% penicillin-streptomycin, 50 U/ml gentamycin (Sigma, St. Louis, MO) and aliquoted into 96-well round-bottom plates (2×10^4 cells/well in a total volume of 100 μ l) in the presence or absence of *L. major* promastigotes at a ratio 2:1 (*L. major*: epidermal cells) and incubated at 37°C for 3 days. The infection rate of epidermal cells was determined by incubating LC-enriched epidermal cells with a mixture of acridine orange (5 U/ml) and ethidium bromide (50 U/ml) as previously described [30].

Syngeneic splenic CD4⁺ T cells were added to epidermal cell cultures (2×10^5) to a final volume of 200 μ l/well, and incubation was continued for another 5 days. CD4⁺ T cells were obtained from spleen cell populations by negative selection. Cells were incubated with a cocktail of mAb (J11d, B220, H35-17, M5/114, 33D1) followed by anti-rat IgG-coupled magnetic beads, and then passed through MiniMACS columns (Miltenyi Biotec, Auburn, CA). CD4⁺ T cells were routinely enriched up to 94% as determined by flow cytometry. The enriched CD4⁺ populations were not contaminated by APC because they did not express MHC class II, as determined by flow cytometric analysis (data not shown).

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