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Indoleamine 2,3-dioxygenase (IDO) induced by *Leishmania* infection of human dendritic cells

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Abstract

Dendritic cells (DC) play a pivotal role in regulating immunity, establishing immunologically privileged tissue micro-environments and maintaining homeostasis. It is becoming increasingly clear that one key mechanism that mediates many DC functions is production of the immunomodulatory enzyme indoleamine 2,3-dioxygenase (IDO). For pathogens that cause chronic infection, exploitation of host DCs is a solution to establish and persist within a host. *Leishmania* parasites cause a range of clinical manifestations, all involving chronic infection, and are proficient at avoiding immune responses. We demonstrate here that infection of human myeloid-derived DC with *L. major* and *L. donovani* induces IDO expression using a mechanism that involves autocrine or paracrine stimulation with a DC secreted factor. *Leishmania*-induced IDO suppresses allogeneic and tetanus toxoid-specific lymphocyte proliferation; an inhibition that is reversed with the IDO inhibitor, 1-Methyl Tryptophan (1-MT). Furthermore, IDO expression by human DC does not require live *Leishmania* infection, as parasite lysates also up-regulate IDO mRNA production. Our data suggest that one mechanism *Leishmania* parasites utilize to circumvent immune clearance may be to promote the induction of IDO among host DC within the infection micro-environment.

Keywords

Leishmania; Indoleamine 2; 3-dioxygenase; dendritic cell

Introduction

Chronicity is a hallmark of parasitic infections. As these pathogens often have complex lifecycles involving multiple host species, they elicit a variety of different immune-evasion strategies to persist, increasing the probability of transmission to their vector hosts. All the major protozoan and metazoan pathogens associated with human disease have evolved multiple sophisticated strategies to avoid immune destruction (1, 2); complexities that, to date, have placed efficacious vaccines outside the reach of development. *Leishmania* parasites employ a variety of different mechanisms to establish and maintain infection in

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their vertebrate hosts (3, 4), and despite substantial efforts, an effective vaccine remains elusive.

Leishmaniasis is endemic in 88 countries with approximately 12 million individuals infected and 350 million more at risk (5). A spectrum of disease manifestations is associated with *Leishmania* infection, ranging from localized cutaneous lesions to fatal visceral disease. Although host genetics are no doubt important, these varied clinical outcomes generally are attributed to differences in the *Leishmania* species initiating the infections (6–9). For example, the old world parasite *L. major* induces cutaneous lesions that heal spontaneously; resolution of this disease usually corresponds with immunity, however, substantial scarring occurs along with recurrent infections in immunocompromised individuals (10). The most debilitating form of infection is visceral leishmaniasis, or kala-azar. This disease is most often associated with infection with *L. donovani* and *L. infantum*. Visceral leishmaniasis is a severe systemic disease, characterized by a lack of cutaneous pathology, parasite dissemination, and uncontrolled growth in the liver, spleen, and bone-marrow. If the symptoms are left untreated, this disease is generally fatal.

Leishmania parasites are notably adept at circumventing immunity. These parasites reside within cells of the mononuclear lineage, primarily macrophages (MP) and dendritic cells (DC). Intracellular survival in these cells requires these organisms be able to escape the host cell anti-microbicidal mechanisms. *Leishmania* avoid intracellular elimination in MP by failing to elicit an oxidative burst upon entry (11, 12), delaying phagosome maturation (13), and inhibiting IFN- γ signaling in infected MP (3). While these pathogens enter naïve MP silently, they are destroyed by IFN- γ primed MP in an NO-dependent manner (14), necessitating that they also suppress or delay the production of activating cytokines.

Leishmania primarily reside in MP, although several cell types can harbor these pathogens including DC, neutrophils, monocytes and even fibroblasts. Given the central role of DC in regulating immune responses, exploiting these cells for immune-mediation provides an evolutionary survival strategy to suppress host cell-mediated immunity. One mechanism by which DC exert their regulatory ability is through regulation of the tryptophan catabolism pathway wherein indoleamine 2,3-dioxygenase (IDO) converts tryptophan to kynurenine (15). IDO-producing DC deplete local tryptophan levels during polyclonal T-cell responses (16), supporting the idea that IDO-producing DC inhibit T-cell proliferation. Once T-cell proliferation has been halted by a lack of tryptophan, T-cell death is promoted through exposure of anergic T-cells to toxic kynurenines, exerting differential effects on the Th1/Th2 balance (17–20) and skewing the balance toward a Th2 bias.

The importance of IDO during infectious disease events is only beginning to emerge. Previous studies on *Toxoplasma* spp. and β -Streptococci revealed that IDO is able to deprive an invading organism of tryptophan, thereby halting their replication (21–23). It has also been shown that IDO is induced as a result of infection with HIV (24), influenza (25), hepatitis C (26), and *Plasmodium yoelii* (27). More recently, it has been reported that *L. major* induced IDO expression reduces T-cell proliferation and attenuates pro-inflammatory responses in a murine model system (28). Here, we demonstrate that *Leishmania* infection induces IDO production in human DC and that inhibition of IDO restores the ability of infected DC to activate a proliferative response. Our results suggest that IDO induction is involved in the immunosuppression associated with *Leishmania* infection in humans.

Materials and Methods

Dendritic Cell Generation

Human monocytes (CD14⁺) were isolated from peripheral blood samples of healthy adult donors (Indiana Blood Distribution) using the Ficoll-Paque (MediaTech) method followed by selection using anti-CD14 magnetic microbeads (Miltenyi). The studies described herein were exempt from Ethics Board Review, according to US National Institutes of Health Guidelines (Exemption 45 CFR § 46.101(b)(4)). DC were differentiated from monocytes by IL-4 and GMCSF supplementation, as previously described (29). For mixed lymphocyte reactions (MLR), the same differentiation procedure was utilized except that the media used was serum-free X-vivo media (BioWhittaker).

Parasite Infection Assays

Leishmania major NIH Friedlin V1 strain and (MHOM/IL/80/FN), *Leishmania donovani* 1S strain (MHOM/SD/62/1S), were cultured in 1X M199 media supplemented with 20% FBS, 100mg/mL Penicillin/Streptomycin, and 2mM L-glutamine, and incubated at 26°C. Metacyclic promastigotes were isolated as previously described (30). *Leishmania* spp. were complement opsonized with 5% normal human serum before infecting at a concentration of 10 parasites: 1 DC; infection rates were determined by Diff-Quick staining and were between 63 and 82% with infections between species not differing by more than 12%. For transwell experiments, equal numbers of DC were plated in (lower) wells and (upper) 0.4 micron inserts (Becton Dickinson Laboratories). Lysates were generated by three successive and rapid freeze-thaw cycles (-80°C and 37°C) with brief waterbath sonication of parasite samples equivalent to the 10:1 infection concentrations noted earlier.

Plasmodium falciparum strain 3D7 was maintained as described by Trager and Jensen (31). *Schizonts* were magnetically selected prior to exposure to human DC. Briefly, infected red blood cell (RBC) cultures were washed in RPMI-1640 and applied to an LS column (Miltenyi) in a Vario Macs magnet. The column was washed, removed from the magnet, and mature schizonts were eluted.

Proliferation Assays

For MLR, 2.0×10^4 non-adherent DC mixed with 4.0×10^5 allogeneic lymphocytes (CD14⁻ fraction from monocyte separation) and *Leishmania* (10 parasites:1 DC) were cultured for 5 days in a tryptophan defined media (32). 1-methyl tryptophan (1-MT) (Sigma-Aldrich) was added to wells at day 0 at a concentration of 200µM, as previously reported (16) to inhibit IDO activity. Proliferation was assessed by the addition of 1µCi/ml ³H- thymidine (Perkin Elmer) the last 18 hrs of culture. Antigen-specific proliferation was assessed as described for MLR, except DC were pulsed with 100 µg/ml tetanus toxoid (Astarte Biologicals) and autologous responder cells were utilized. Statistical significance was determined using a Student's T-Test with a confidence level of 95%.

Quantitative RT-PCR and Analysis

RNA was isolated using an RNeasy Mini kit (Qiagen) and cDNA was generated using SuperScript III (Invitrogen) following manufacturer protocols. Quantitative real-time PCR (qRT-PCR) reactions were performed using SYBR Green Mastermix (Applied Biosystems) was utilized as previously described (33), with 25ng cDNA and 300nM Forward and Reverse primers (IDT) per reaction. Human primers for *HPRT1*, *IDO1*, *IFNA1*, *IFNB1*, and *IFNG* mRNA transcripts were used (see Supplemental Table 1 for primer sequences). Fold change over uninfected gene expression levels were calculated using the delta delta Ct method normalized to *HPRT*. Significance levels were determined with the Mann-Whitney

test or one-way repeated measures ANOVA with Bonferroni post-tests using GraphPad Prism 5.0 software.

Flow Cytometry

Samples fixed with 2% paraformaldehyde were washed 3 times with 0.1% BSA – PBS. Cells were blocked in PBS supplemented with normal mouse serum (Jackson Immunoresearch) and BSA for 1hr and subsequently stained with a panel of antibodies: isotypes, anti-CD40-PE (5C3), anti-CD80-PE (L307.4), anti-CD83-PE (HB15e), anti-CD86-PE (FUN-1), and anti-HLA-DR-PE (L243) (BD Pharmingen). All samples were washed and then analyzed using an MCL500 flow cytometer (Beckman Coulter). Data was expressed relative to uninfected controls.

Western Blot Analysis

DC were lysedfor 20 min with ice-cold lysis buffer (34). Proteins were separated by SDS-PAGE and transferred to nitrocellulose membranes. Blots were probed with primary antibodies against IDO (Chemicon) and GAPDH (Biogenesis), incubated with a secondaryHRP-conjugated anti-mouse Ig antibody (BD Biosciences)in PBST plus 5% BSA. Antibodies were detected usingSuperSignal West Pico and Femto ECL reagents (Pierce).

Results and Discussion

IDO Expression in Human Dendritic Cells

Our initial observations of IDO expression in *Leishmania* infected DC were made by remining previously published microarray data (35). These microarray data represent pooled monocyte-derived DC generated from blood samples of seven healthy adult human donors that were subsequently infected *in vitro* with either *L. major* or *L. donovani*. RNA was harvested 16 hours post-infection, pooled, and hybridized to an Affymetrix HU95A microarray (35). Our data analysis revealed that IDO is induced in human DC 10.35 and 1.55 fold in response to *L. major* and *L. donovani* infection respectively compared with uninfected DC (data not shown).

In order to confirm these results, human monocyte-derived DC from six additional healthy human blood donors were infected in vitro with either L. major or L. donovani. Following 16 hours of infection, RNA was harvested, and quantitative RT-PCR (qRT-PCR) was utilized to measure IDO expression from each donor individually. IDO mRNA was significantly induced by infection with *L. major* or *L. donovani* (p 0.05, Fig. 1 A). Although not statistically significant, the levels induced by L. donovani were generally lower than that induced by L. major. DC from another six donors were utilized to determine the translational state of IDO during the infections detailed above (Fig. 1B). IDO protein also was induced by Leishmania infection, and similar to IDO mRNA expression, expressed at a lower level in L. donovani-infected DC as compared to L. major infected DC. To determine if IDO induction requires live Leishmania infection we stimulated DC with L. major lysate (Fig. 1C). Parasite lysates induced significant IDO mRNA expression, though to a lower level than during live L. major infections. To distinguish between the effect of phagocytosis versus infection with Leishmania parasites, we exposed DC to uninfected and P. falciparum-infected allogeneic RBC and assessed the induction of IDO by qRT-PCR (Suppl. Fig. 1A). IDO expression was not induced by uninfected RBC uptake, suggesting that the effect observed in response to *Leishmania* infection is not solely due to phagocytosis. Interestingly, IDO mRNA is up-regulated in response to P. falciparuminfected RBC in human DC, a response that results in detectable protein (data not shown).

An early pro-IDO signal is likely mediated through pathogen interaction with pattern recognition receptors, such as toll like receptors (TLR) on the DC surface, as many TLR have been shown to induce IDO (36). Interestingly, Leishmania species have been shown to interact with TLR2 and TLR3 (37, 38) and effective control of L. major infection in vivo is partially dependent on TLR4 (39). IDO also is induced by cytokines, in particular interferons (40). To investigate if autocrine or paracrine regulation is involved in Leishmania induction of IDO, we utilized a transwell system whereby infected DC in the upper chambers could produce soluble mediators to affect IDO production in uninfected DC in the lower chambers (Fig. 1D). We infected cells in the lower chamber with L. major parasites, where these cells expressed IDO as expected. This induction appears to be mediated partially by a soluble factor(s) as IDO expression was up-regulated in uninfected DC exposed only to infection via the physically segregated upper transwell. Infection in both chambers, however, resulted in synergistic induction of IDO. Two possible explanations can account for this latter result. First, increased IDO expression may only result from the additive effect of more soluble mediators being present due to increased L. *major* infection (*L. major* in both wells). Alternatively, the increased expression may be the result of two signals, one that requires direct contact and one resulting from an autocrine mechanism.

IDO Transcript Expression Follows a Model of Interferon Expression Over Time

To assess gene transcript regulation during the early infection period, we used qRT-PCR to monitor the expression of IDO in DC from five healthy adult human blood donors at 2, 4, 8, and 24 hours post-*L. major* infection (Fig. 2A). The fold induction of IDO among infected DC compared with that of uninfected DC increased significantly after 8 hours and continued through to 24 hours. These data agree with our earlier qRT-PCR observations at 16 hours post-infection and suggest a progressive IDO induction as a function of *Leishmania* presence and engagement of host cells, within the cellular micro-environment.

The *INDO* (gene encoding IDO) promoter contains several interferon response elements (40), suggesting that the soluble mediator(s) described in either of the earlier explanations for the transwell assay results could be an interferon. Of interest is that L. major infection activates interferon regulator factors 1 and 8, as well as STAT1 in human DC-three factors that are able to bind and activate interferon response elements (41). qRT-PCR was again utilized with DC samples of the five healthy blood donors used in time course infection assays tracking expression of the three primary types of interferons (IFNa, β , and γ). IFNa mRNA levels displayed a modest up-regulatory trend over time (Fig. 2B), achieving significance compared with uninfected samples only after 24 hours post-infection. Relatively lower levels of IFNa mRNA suggests this interferon is not responsible for IDO induction, as would be expected considering the pathological role IFNa has been shown to play in aberrant IDO promotion (42). Like IDO, IFN γ was significantly up-regulated 8 hours post-infection. However, unlike IDO, IFN γ transcript levels began to decrease at 24 hours post-infection (Fig. 2D). Interestingly, IDO expression levels do not drop at 24 hours post-L. major infection, but instead continue to rise after IFNy begins to fall (Fig. 2A). IFN β expression peaked at approximately 4 hours post-infection, but progressively dropped back to insignificant levels by 24 hours post-infection (Fig. 2C). There is evidence supporting a role of IFN β positively influencing secretion of autocrine-acting IFN γ by DCs (43); and it has also been shown that IFN γR engagement stimulates the translocation of IRF1 and IRF8 (ICSBP) to the DC nucleus, promoting immune molecules' expression in a conditional manner (44). In concordance with these reports, the interferon expression profiles following L. major infection suggest a possible stepwise model of earlier IFNB expression, its subsequent promotion of IFN γ expression followed by IDO induction.

Alternatively, IFN β may directly mediate IDO up-regulation as has been reported for human trophoblast cells (45).

Infection-Induced IDO is Active and Inhibits a Mixed Lymphocyte Reaction

In some systems IDO mRNA and protein is produced but the protein is not functional (16, 46); to ensure that IDO induced by *Leishmania* infection (Fig. 1B) was functional, mixed lymphocyte reactions (MLR) were utilized. Active IDO inhibits T-cell polyclonal activation associated with MLR, and the addition of 1-Methyl Tryptophan (1-MT) reverses this inhibition (47). Our MLR results indicated that IDO is active as conditions that include *Leishmania* infection exhibit significantly reduced levels of proliferation relative to uninfected allogeneic conditions (Fig. 3). This infection-induced inhibition is reversed by the addition of the IDO inhibitor, 1-MT. To further demonstrate that IDO induced by *L. donovani* infection is active, we supplemented the MLR media with L-tryptophan (Suppl. Fig. 2). Addition of exogenous tryptophan significantly reversed the proliferation inhibition induced by *L. donovani* infection of DC.

To assess if *Leishmania*-infection induced IDO also is active against antigen-specific cell activation, a tetanus toxoid-based proliferation assay was utilized. DC from healthy human blood monocyte donors were infected and co-incubated with autologous responder cells with or without 1-MT. Patterns in antigen-specific proliferation were similar to those observed with MLR, as infection with either *L. major* or *L. donovani* inhibits proliferation and the addition of 1-MT circumvents the infection-based inhibition (Fig. 4). Conversely, the IDO induced in response to uptake of *P. falciparum*-infected RBC does not inhibit antigeneic-specific cell proliferation (Suppl. Fig. 1B).

Even though previous studies have demonstrated that *Leishmania* infection has little impact on co-stimulatory markers (29, 48), to rule out the role of decreased surface expression in inhibition of proliferation, surface level expression of CD40, CD80, CD86, and HLA-DR was assessed by flow cytometric analysis (Suppl. Fig. 3). No significant differences were observed in *Leishmania*-infected DC compared to uninfected controls. While previous literature demonstrates a connection between immunosuppression and *Leishmania* infection (49, 50), no direct mechanism influencing this phenomenon has been defined in humans. Our studies imply a role for IDO in immunosuppression associated with leishmaniasis.

A recent murine study demonstrates that IDO contributes to the *L. major* parasite burden and footpad swelling indicating that IDO expression is advantageous for pathogen survival (28). A few novel studies have investigated the potential of *Leishmania* infection induced immunosuppression to alter a host's response to other organisms or vaccines. In BALB/c mice, *L. major* infection ablates the cellular mediated response induced by a previous vaccination against the HIV-1 gag gene (51). In a clinical case report, an individual presented with classical co-infections associated with HIV (i.e., tuberculosis, pneumoncystosis, and strongyloidiasis). Those infections persisted despite their specific treatments and finding that the patient was HIV negative. However, he was later found to be infected with *L. infantum chagasi* as a co-morbidity. Treatment with antimony-based compounds led to the patient making a full recovery, with clearance of all other pathogens after the *L. i. chagasi*-induced immunosuppression was alleviated (52). No study has provided a definitive mechanism for this noted *Leishmania*-induced immunosuppression, although the interaction of CTLA-4 with B7 (53) has been implicated. Interestingly, this same interaction is a potent inducer of IDO (16).

Concluding Remarks

Our report demonstrates that active IDO is induced in human DC in response to infection with *L. major* or *L. donovani*, leading to inhibition of both polyclonal and antigen-specific proliferative responses. Together, these data indicate that one of the mechanisms behind the establishment of *Leishmania* spp. infection is through the induction of DC-expressed IDO.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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ABBREVIATIONS

IDO	Indoleamine 2,3-dioxygenase

1-MT 1-Methyl Tryptophan

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Donovan et al.

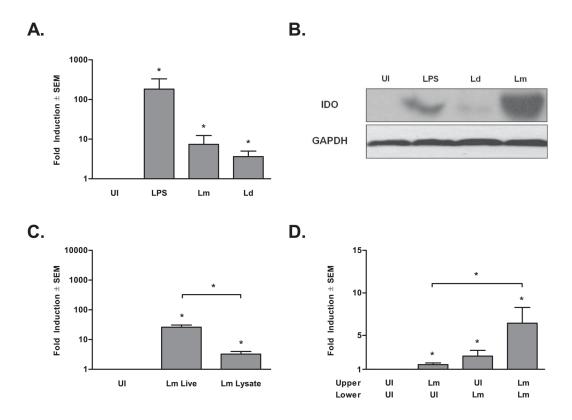


Figure 1. IDO is Induced in Human DC in Response to Leishmania Infection

A)*IDO* mRNA transcript expression as measured by qRT-PCR using equal concentrations of cDNA from six healthy adult human monocyte-derived DC and normalized to *HPRT*. Cells were left uninfected (UI) or infected *in vitro* with *L. major* (Lm) or *L. donovani* (Ld) or exposed to 1µg/mL of LPS (LPS) for 16 hrs. B) Western blot analysis of human monocyte-derived DC infected with *L. major*, *L. donovani*, or stimulated with LPS (1µg/mL) for 48 hrs. Blots were probed for IDO and GAPDH as a loading control. Data from one of six representative donors is presented. C) qRT-PCR of *IDO* mRNA from human monocyte-derived DC, infected *in vitro* with live *L. major* or stimulated with an equivalent amount of parasite lysate for 16 hrs. D) *IDO* mRNA detected from uninfected or *L. major* infected DC from the lower chamber of a transwell system. Infection in upper wells as indicated, n = nine donors. Data is expressed as mean \pm standard error of th mean (SEM) fold induction over uninfected controls, *p 0.05 by Mann-Whitney Test.

Donovan et al.

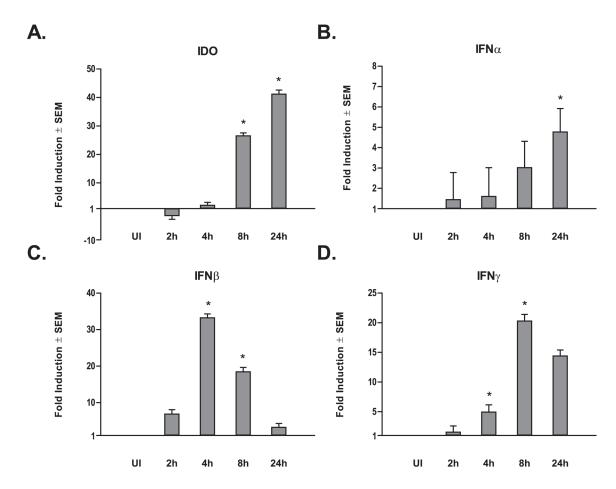


Figure 2. *Leishmania* Infection Affects Differential Indoleamine and Interferon Transcript Expression Patterns over Time

qRT-PCR for A) *IDO*, B) *IFNa*, C) *IFN* β ,and D) *IFN* γ normalized to *HPRT* transcript expression levels using equal concentrations of cDNA from five healthy adult human monocyte-derived DC. Cells were either uninfected or infected *in vitro* with *L. major* and samples extracted after 2, 4, 8, and 24 hours post-infection. Data is expressed as mean transcript expression fold change over uninfected controls normalized to *HPRT* ± SEM. *p

0.05 compared with uninfected (UI) gene expression levels by one-way repeated measures ANOVA and Bonferroni multiple comparisons tests.

Donovan et al.

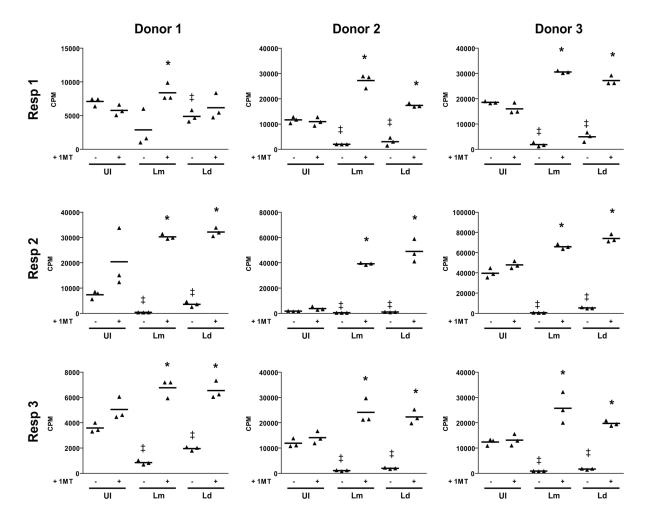


Figure 3. Leishmania-Infection Induced IDO is Active and Inhibits MLR

DC from nine healthy donors were either left uninfected (UI) or infected with either *L. major* (Lm) or *L. donovani* (Ld) and then mixed in auto- and allogeneic conditions at a ratio of 20 responders to 1 DC and incubated for 5 days. Proliferation was assessed by ³Hthymidine incorporation the last 18 hrs of culture. Data is expressed as β -particle counts per minute (CPM) with conditions listed with or without 1-MT. [‡] p 0.05 compared to uninfected control; * p 0.05 compared to same infection lacking 1-MT as determined by Student's T-Test. Data from three representative donors out of nine are shown.

Donovan et al.

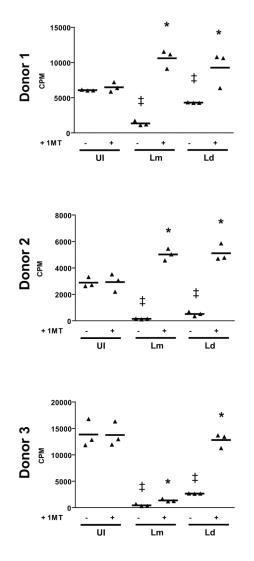


Figure 4. *Leishmania*-Infection Induced IDO is Active and Inhibits Antigen-Specific Proliferation

Uninfected DC (UI) or DC infected with either *L. major* (Lm) or *L. donovani* (Ld), autologous responders, and 0.10 ng/mL tetanus toxoid were cultured for 5 days. Proliferation was assessed by ³H-thymidine incorporation the last 18 hrs of culture. Data is expressed as counts per minute (CPM) with conditions listed with or without 1-MT. [‡] p 0.05 compared to uninfected control; * p 0.05 compared to same infection lacking 1-MT as determined by Student's T Test.