

IMMUNOPATHOLOGY OF LEISHMANIASIS: AN UPDATE

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Leishmaniasis represents a severe, increasing, public health problem. The perspective of its control is highly dependent on research progress, on therapeutic manipulations of the immune system, and on vaccine development. There is a correlation between the clinical outcome of *Leishmania* infection and the cytokine response profile. While a protective immune response against *Leishmania* has been clearly identified to be related to the influence of a type-1 response and IFN- γ production, the precise role of T helper (T_H) 2 cytokines in non-healing infections requires further exploration. IL-4 and IL-13 (T_H2 cytokines) can promote disease progression in cutaneous leishmaniasis, whereas IL-4 would appear to enhance protective type-1 responses in visceral leishmaniasis. Thus, the T_H1/T_H2 paradigm of resistance/susceptibility to intracellular parasites is probably an oversimplification of a more complicated network of regulatory/counter regulatory interactions. Moreover, the presence of antigen specific regulatory T cell subsets may provide an environment that contributes to the balance between T_H1 and T_H2 cells. Finally, the involvement of CD8⁺ T cells has been described, but the modality of their function in this kind of infection has not been so far elucidated.

Leishmaniasis represents a severe, increasing public health problem (1-2). Moreover, many of the 50 million people who travel from industrialized to developing countries each year report some infective illness associated with their travel, included leishmaniasis, which is endemic in the areas of tropics, subtropics, and southern Europe (3). The perspective of control is still highly dependent on research progresses, to obtain better tools and more cost-effective strategy for vector control and case management, through therapeutic manipulations of the immune system, and vaccine development (4).

Leishmaniasis comprises a group of diseases caused by protozoan parasites of the *Leishmania* genus. Leishmaniasis is characterized by different clinical manifestations, which are dependent not

only on host genetic control, but also on the infecting species, which to date are more than 20 (5).

The parasites which cause the various forms of leishmaniasis in humans are classified in the subgenus *Leishmania* (*L.*) or *Viannia* (*V.*). Cutaneous leishmaniasis (CL) is caused, in the Old World, by *Leishmania* (*L.*) *major*, *Leishmania* (*L.*) *aethiopica*, and dermatropic *Leishmania* (*L.*) *infantum*, of the *Leishmania* subgenus; in the New World, by *Leishmania* (*L.*) *mexicana*, *Leishmania* (*L.*) *amazonensis*, *Leishmania* (*L.*) *venezuelensis*, and dermatropic *Leishmania* (*L.*) *infantum*, of the *Leishmania* subgenus, and *Leishmania* (*V.*) *braziliensis*, *Leishmania* (*V.*) *guyanensis*, *Leishmania* (*V.*) *lainsoni*, *Leishmania* (*V.*) *naiffi*, *Leishmania* (*V.*) *panamensis*, *Leishmania* (*V.*)

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peruviana, and *Leishmania shawi*, of the *Viannia* subgenus. Mucocutaneous leishmaniasis (MCL), or espundia, is caused by *Leishmaniasis (V.) braziliensis*, and *Leishmania (V.) panamensis*, whereas visceral leishmaniasis (VL), also known as 'Kala Azar', is caused by *Leishmania (L.) infantum* in the Mediterranean area, Middle-East and Asian countries, and *Leishmania (L.) chagasi* in Latin America. *Leishmania (L.) mexicana* and *Leishmania (L.) amazonensis* are now regarded as a single specie, due to their genotypic relationships, within the *Leishmania mexicana* complex, whereas *Leishmania (L.) infantum* and *Leishmania (L.) chagasi*, together with *Leishmania (L.) donovani*, are now regarded as a single specie, due to their genotypic relationships, within the *Leishmania donovani* complex. After recovery, patients may develop a chronic CL form, called 'Post-Kala Azar Dermal Leishmaniasis' (PKDL), which usually requires long and expensive treatment (6-7).

Leishmania spp. are digenetic parasites, that develop as flagellated motile promastigotes in the gut of blood-sucking female sand flies (*Phlebotomus spp.* in the Old World, *Lutzomyia spp.* in the New World), transmitted into the dermis of the vertebrate host during the ingestion of a blood-meal, and as obligate intracellular non-motile amastigotes in the phagolysosomes of macrophages and dendritic cell lineage of the vertebral hosts. Leishmaniasis is mainly a zoonotic disease and a wide range of mammals act as reservoirs, particularly rodents, edentates and marsupials in CL, wild canines and domestic dogs in zoonotic VL. In anthroponotic VL (widespread in India) man is the sole source of infection for the vector (8-9).

All species of *Leishmania* infect mice relatively easily, producing diseases facilitating their utilisation as *in vivo* models, not only to evaluate the host genetic control of infection, but also to determine how immune responses develop and are regulated (10-11).

Immune responses in animal and human leishmaniasis

General aspects

Parasite factors and host mechanisms are inextricably linked with the pathogenesis of leishmaniasis. To initially establish the infection,

promastigotes, inoculated with the sandfly saliva, enter macrophages to evade host responses. Progressive intracellular infection by amastigotes depends on the maintenance of macrophages in an inert, deactivated state. At the same time, however, the immunocompetent host is equipped to respond both with non-specific (innate) and antigen-specific (adaptive) cell-mediated mechanisms. These inflammatory responses mediate disease evolution and may (self-healing disease or asymptomatic infection) or may not (non-healing disease) produce the desired clinical end-result (8, 12). At the site of infection, innate responses include cells (neutrophils, monocytes, macrophages, natural killer [NK] cells, and dendritic cells), Pattern Recognition Receptors (PPRs, i.e. toll-like receptors), and soluble products (complement, and released cytokines, including interleukin [IL]-1 α , IL-12, Tumor Necrosis Factor [TNF]). Innate mechanisms, especially IL-12 secretion, drive the parallel induction of cell-mediated immunity. This complex set of mechanisms leads to the activation of specific CD4⁺ and CD8⁺ T cells. These effector T cells circulate, are recruited to cutaneous or visceral sites via adhesion molecules and chemokine mechanisms, and, along with influxing blood monocytes, direct local inflammatory responses, including granuloma assembly and lesion development (13-15).

The site of *Leishmania* infection in the dermis of patients affected by CL is characterized by a massive infiltration of macrophages and chemokines, such as Monocyte Chemoattractant Protein-1 (MCP-1), monokine induced by interferon (IFN)- γ (MIG), IFN- γ -inducible protein 10 (IP-10), and only a low amount of macrophage inflammatory protein 1 α (MIP-1 α) (16-17). In skin lesions of patients with self-healing CL, the healing process is facilitated by high levels of MCP-1 expression, whereas high local levels of MIP-1 α are associated with the non-healing form of CL. MCP-1 and IFN- γ synergistically activate monocytes to clear intracellular parasites, whereas IL-4 abrogates the effect of MCP-1 (16-17). Predominant T helper (T_H)1-type cell responses are associated with IFN- γ -induced macrophage activation, indicating a network of pleiotropic cytokines, in which IL-12, produced by activated antigen presenting cells (APC), such as macrophages and dendritic cells,

Table 1. Types of CD4+ T-cells and cytokines involved in different forms of leishmaniasis.

Disease	Healing		Non healing	
	CD4+ T-cell	Cytokines	CD4+ T-cell	Cytokines
Cutaneous leishmaniasis	T _H 1	<i>Leishmania major, mexicana, amazonensis</i> : IL-12, IL-1 α , IL-18, IL-23, IL-27, IFN- γ , TNF- α , MIF, IFN- α , IFN- β , GM-CSF, IL-10, TGF- β	<i>Leishmania major</i> : excessive T _H 2 response and/or defective APC function and/or defective T _H 1 response	IL-4 (?), IL-13, IL-10, TGF- β
			<i>Leishmania mexicana</i> : excessive T _H 2 response and/or defective APC function and/or defective T _H 1 response	IL-4, IL-13 (?), IL-10 (?)
			<i>Leishmania amazonensis</i> : excessive T _H 2 response and/or defective APC function and/or defective T _H 1 response	IL-4 (?)
Visceral leishmaniasis	T _H 1 T _H 2 (?)	IL-4 (?), IL-12 (experimental visceral leishmaniasis)	<i>Experimental visceral leishmaniasis</i> : defective T _H 1 response (?) excessive T _H 2 response (?)	IL-4 (?), IL-10, TGF- β (?)
	T _H 1 T _H 2 (?)	IL-2, IFN- γ , IL-12, IL-15, IL-4 (?) (human visceral leishmaniasis)	<i>Human visceral leishmaniasis</i> : defective T _H 1 response (?) excessive T _H 2 response (?)	IL-4 (?), IL-10, TGF- β (?)

CD: cluster of differentiation T_H1: T helper 1 T_H2: T helper 2 APC: antigen presenting cells IL: interleukin

IFN- α : interferon- α IFN- β : interferon- β IFN- γ : interferon- γ MIF: Migration Inhibitor Factor

TGF- β : Transforming Growth Factor- β TNF- α : Tumor Necrosis Factor- α GM-CSF: Granulocyte Macrophage-Colony Stimulating Factor

shapes the basic response, and IFN- γ and other cytokines also participate (18). However, in patients with clinically apparent infections, especially acute VL (see below), T_H1- and T_H2-type responses are not characteristically polarised, as both activating (i.e. IL-12, IFN- γ) and suppressive (i.e. IL-4, IL-10, IL-13, Transforming Growth Factor [TGF]- β) cytokines are detected (19-20). It is possible that in acute leishmaniasis counterbalancing mechanisms are normally produced to curtail the process (19-20). IL-4, IL-10, IL-13 and TGF- β are capable of derailing T_H1-type responses and deactivating macrophages, thereby moderating tissue injury, but promoting intracellular infection (19-20).

Resolution of infection: the T_H1 response

It is now well established that a protective immune response against CL is dependent on the development of a potent type-1 response (15). The general consensus is that IL-12, from activated APC, possibly augmented by other cytokines (IL-1 α , IL-18, IL-23 and IL-27, as innate mechanism) drives the differentiation and proliferation of T_H1 cells, which

produce, together with NK cells, IFN- γ , as antigen-specific acquired cell-mediated response (21-23).

Major Histocompatibility Complex (MCH) class II antigen presentation alone is not sufficient to activate T_H1 cell responses and ligation of co-stimulatory molecules (i.e. B7-1/B7-2 and CD40 on the APC with CD28 and CD40L on the T_H1 cell, respectively) is also a prerequisite (24-25). IFN- γ from T_H1 cells, and probably to a lesser extent CD8⁺ T cells (as part of the antigen-specific acquired cell-mediated immune response), but also from IL-12 activated NK cells, as part of the innate response, mediate macrophage activation, superoxide (O₂⁻) and nitric oxide (NO) production, and parasite killing. Although the molecular mechanism of the action of NO on *Leishmania* is to date unknown, early data suggested that NO is directly cytotoxic to the parasites (26-27). The NO-pathway thus appears to be a common mechanism of *Leishmania* killing. Moreover, macrophage leishmanicidal activity, induced by IFN- γ , has been shown to be enhanced by other cytokines, such as TNF- α , Migration Inhibitor Factor (MIF) and type-1 interferons

(IFN- α subtypes and IFN- β subtype), Granulocyte Macrophage-Colony Stimulating Factor (GM-CSF), as well as by CD40/CD40L interactions (28-29).

For the development of an adequate T_H1 -mediated resistance to *Leishmania* infection, another important pathway is the Fas/FasL interaction. Activated T_H1 cells induce apoptotic death in target cells expressing the Fas protein (30). Macrophages, infected with *Leishmania*, upregulates Fas, in response to IFN- γ , and become susceptible to CD4⁺ T cell-induced apoptotic death (30-31). Thus, Fas-induced apoptotic death of infected macrophages might limit the number of host cells at the site of infection which are required for amastigote replication (30-31).

Following resolution of infection, concomitant immunity is dependent upon antigen-specific CD4⁺/CD25⁺ T-regulatory cells, producing IL-10 and TGF- β , that moderate the activity of T effectors (32-33). In the absence of persistent infection, protective immunity is significantly reduced (32-33).

Non-healing cutaneous Leishmania major infection: T_H2 -dependent or defective T_H1 response

The T_H1 - T_H2 paradigm of resistance/susceptibility to intracellular infection is largely based on investigations using *Leishmania major*. Initial studies have suggested that the resolution or progression of disease is dependent on distinct CD4⁺ T cell subsets, T_H1 and T_H2 , producing the counter regulatory cytokines IFN- γ and IL-4, respectively (34-35). However, contradictory reports have been published indicating a disease progression role for IL-4 (36). These studies posed two significant questions: firstly, could other regulatory cytokines be responsible for the immunosuppressive activity previously attributed to IL-4, and/or secondly, could the well documented defective T_H1 response play a major role in progressive disease?

Recent studies, using gene-deficient and transgenic mice, have clearly identified other cytokines in addition to IL-4 having major roles in the non-healing response to *Leishmania major* (36-37). Particularly, IL-13 has been found to act independently of IL-4, and the effects of IL-13 and IL-4 might be additive (37). Other studies suggest that IL-10 is at least as influential as IL-13 and IL-4 in promoting *Leishmania major* disease progression.

IL-10 might play a key role in the pathogenesis of leishmaniasis, especially in the down-regulation of T_H1 responses, as supported by the observations of suppression of IL-12 production by APC, and of IFN- γ production by CD4⁺ T and NK cells, and of macrophage deactivation and inhibition of their leishmanicidal functions. T regulatory cells are also significant producers of TGF- β , which is also partly responsible for suppressing protective responses (36-38).

The inability to mount a T_H1 response, irrespective of a T_H2 response, has been attributed to an inability to produce or respond to IL-12, and, therefore, to intrinsic defects in APC function or in T_H1 cell development. Defective APC function, attributable to a failure to produce IL-12, may also underlie in a deficient APC IL-1 production (39). As a matter of fact, IL-1 upregulates IL-12 production, as well as MHC class II co-stimulatory molecule expression, and mediates the ability of IFN- γ to inhibit T_H2 response (39).

On the contrary, intrinsic defective T_H1 cell development, originally identified as an inability to respond to IL-12, has been associated with a downregulation of T_H1 cell IL-12R β 2 expression (40). It is well established that IL-12R β 2 expression can be downregulated by IL-4 produced by T_H2 , even though IL-4-independent mechanisms have also been described (40). A recently described defective co-polarization of the T cell receptor (TCR) and the IFN- γ receptor complex in naive CD4⁺ T cells during the APC/T precursor interaction, would also significantly favour a commitment to T_H2 development (39-40).

Non-healing cutaneous Leishmania mexicana/Leishmania amazonensis infection: IL-4-dependent

In non-healing cutaneous *Leishmania mexicana* infection IL-4 appears to exert the major contribution, whereas IL-13 and IL-10 are only partially involved (41). The Cathepsin L-like cysteine peptidase (CPB) has been identified as the likely *Leishmania mexicana/Leishmania amazonensis* virulence factor inducing IL-4 production (42).

While studies on *Leishmania mexicana* have clearly shown a major disease exacerbating role for IL-4, that might subvert the development of a T_H1 -response, similar studies, on the closely related

parasite *Leishmania amazonensis*, have suggested an insignificant role for IL-4 in non-healing infection (43-44). In this setting, CPB might be directly responsible for the inhibition of T_H1 responses, due to the ability of CPB to proteolytically degrade the NF- κ B family of signalling proteins (43-44). Thus, IL-4 independent mechanisms might prevail (43-44).

Paradox of T_H1 and T_H2 in visceral leishmaniasis

The dichotomy of T_H1 and T_H2 cell responses induced by *Leishmania* parasites is even less clear in VL. In experimental VL by *Leishmania donovani*, the disease progression is due to the failure of an appropriate T_H1 response, rather than to the T_H2 cell proliferation (45). In fact, IL-4 has been shown to have no exacerbating role in murine VL, and, under certain circumstances, it can prime for IL-12 production and a type-1 response (41).

In humans infected with *Leishmania donovani* high concentrations of IL-10 and IFN- γ are detected in the sera at the beginning of infection, which return to the normal range following successful chemotherapy (46-47). By contrast, peripheral blood mononuclear cells (PBMC), stimulated *in vitro* with *Leishmania* antigen, produce low levels of IL-10 and IFN- γ when collected at the time of the diagnosis, and normal levels when assayed after recovery (46-47). IL-4 and IL-2 are undetectable in the sera, whereas IL-4 is in high concentrations in the supernatants in all the phases of the disease (46). IL-2 is significantly reduced in the supernatants of actively infected patients, returning to the normal level after recovery (46-47). Low sera levels of IL-4 might be related to the high production of IFN- γ , whereas reduction in IL-2 sera levels might be due to the high sera concentration of soluble IL-2 receptor (sIL-2R) that binds IL-2 with a high affinity mechanism (48). In spite of the similar secretion pattern of IL-10 and IFN- γ , above reported, the control of infection or complete recovery seems to be associated with an increased production of IL-2 and IFN- γ (49-51). Furthermore, IL-10 production correlates with the progression of VL (52), and neutralisation of IL-10 with a specific monoclonal antibody restores T cell proliferation and IFN- γ production in PBMC from acute VL patients (53). *In vitro* studies have also demonstrated that

IL-12 shifts the responses toward a T_H1 -type and enhances IFN- γ production (54). These findings therefore suggest that, in patients with active VL, the cytokine profile is not clearly polarized, and both T_H1 -like and T_H2 -like cells appear to proliferate and to be activated (46, 50, 55). The balance of cytokines at the site of primary activation of the *Leishmania*-specific cells appears to be of major importance for the development of T_H1 and T_H2 responses (56-57), even though other unknown factors might influence the cellular immune response. Altogether, data indicate that in symptomatic patients T_H1 cytokine production is not depressed, but there is an unresponsiveness to the stimuli of these cytokines, also caused by the high production of IL-10. Recently, another lymphokine, IL-15, was studied for its role against infectious diseases (58) and for its ability to enhance both T_H1 responses, by increasing IFN- γ production from NK and T cells (59), and T_H2 responses, by augmenting IL-5 and IL-13 production (60). In particular, it has been demonstrated that endogenous IL-15 plays a role in the suppression of T_H2 cytokines in acute VL patients, even though it does not enhance the production of T_H1 cytokines (61), indicating a potential protective role of IL-15 against leishmaniasis caused by the indirect effect on T_H1 , due to the restriction of T_H2 proliferation (61). Furthermore, it has been demonstrated that the activation of macrophage cultures with IL-15 determines a significant anti-*Leishmania* activity (62). The killing of *Leishmania* in macrophages, primed with IL-15, is followed by an increase in the IL-12 synthesis (62). Altogether, these data indicate that IL-15 could have a role as an activator of leishmanicidal activity, directly or indirectly, by inducing IL-12 production.

Non-healing visceral leishmaniasis: T_H2 -independent?

As already mentioned, the disease progression and the exacerbating role for T_H2 response and IL-4 production during *Leishmania donovani* infection has yet to be demonstrated. Early studies suggest that the differential production of T_H1 and T_H2 cytokines does not control the rate of cure (45), although the production of IFN- γ correlates with resistance, T_H2 cytokines are not so clearly responsible for susceptibility (63).

According to this point of view, studies in mice have shown that animals deficient in IL-4 and IL-4R α are more susceptible to disease than their wild-type counterparts, suggesting that IL-4 may be protective in some circumstances, and can promote resistance, rather than susceptibility, to intracellular pathogens (64). In addition, IL-4 and IL-4R α signalling are not only essential for optimal clearance of *Leishmania donovani* from the liver and for limiting infection in the spleen (following primary infection), but also are effective for T cell dependent chemotherapy, and for vaccine-induced resistance (65). In the absence of IL-4, type-1 responses and IFN- γ production fail to be maintained following chemotherapy or fail to be induced by vaccination (64-65). Other studies indicate that IL-10 is the major immunosuppressive cytokine in VL, although TGF- β has also a significant disease-promoting activity (66). Experimental IL-10 inhibition allows activation of T_H1 cell responses and promotes parasite killing (66-67). Furthermore, while the healing process in susceptible mice is IL-12 dependent, IL-10, but not IL-4 or TGF- β , appears to blunt T_H1-type responses and to determine fatal outcome of the disease (21).

T cell subset modifications in visceral leishmaniasis

Although data indicate that *Leishmania donovani* parasites cause alterations of the immune system, with immune-depression, the exact mechanism by which the parasites induce immune-depression is not clear. In patients with symptomatic *Leishmania infantum* infection the levels of CD3⁺, CD4⁺ and CD8⁺ T cells are within the normal range (46, 50), whereas acute VL patients have markedly reduced levels of memory T cells (CD3⁺/CD45RO⁺) compared with healthy controls, and these cells returned to the normal levels following successful chemotherapy (46, 50, 68-69). On the contrary, the number of memory CD4⁺ T cells is markedly increased in persons with asymptomatic infection and positive leishmanin (or Montenegro) skin test, intradermally performed using a culture of promastigotes as antigen (68). The significant increase of memory CD4⁺ T cells in this group of patients, their relationship with the positive leishmanin skin test and acquired immunity needs further explanations (68-69). However, T cells are strongly activated in acute VL patients, as indicated

by the elevated number of CD3⁺ HLA-DR⁺ and by the increase in HLA-DR antigen on these cells (46).

The role of apoptosis and CD8⁺ T cells in Leishmania infection

Previously, it has been shown, both in mice and in humans, that the induction of T cell apoptosis could be involved in the defective host-cellular responses to challenge with pathogenic infectious agents (70-73). In experimental VL it has been demonstrated that the infection of a susceptible host results in CD4⁺ T cell apoptosis and a decrease in T_H1 cytokine production (73). In particular, it has been demonstrated that resistance to *Leishmania major* depends on apoptotic mechanisms, mainly operating through the Fas (APO-1/CD95) pathway, and singeneic *gld* and *lpr* mice, lacking a functional Fas system, fail to heal their lesions (74). Furthermore, in mice infected with *Leishmania donovani* parasites an increased incidence of T cell apoptosis in liver and spleen was observed (75). Also in VL patients, both in acute and in healed phases of disease, leukocyte apoptosis seems to be involved. In fact, monocytes and T lymphocytes from acute VL patients show a significantly higher level of apoptosis compared with that observed in healed subjects. The percentage of apoptotic cells was higher in monocytes than in T lymphocytes (31). T cells involved in programmed cell death were mainly of CD4⁺ phenotype (31, 76). In particular, the subset T_H1, evaluated by chemokine receptor-5 (CCR5), was involved in this process and used CD95-mediated mechanism (31). The significant increase in apoptosis of T_H1-like subset (15-20% of CD4⁺ cells) is in line with the studies carried out in susceptible hosts, showing an enhancement of CD4⁺ T cell apoptosis, particularly of T_H1-like cells, associated with a decrease in T_H1 cytokine production (73). Since, T_H1 cytokines appear to be involved in the protection against leishmaniasis, both in experimental models (27, 73) and in human infections (57-59), this deletion of CD4⁺ T_H1-like cells could contribute to the depressed cell-mediated immunity in acute VL patients.

On the contrary, in active human localised CL a higher frequency of cell death in CD8⁺ T cells than in CD4⁺ T cells was detected, and these apoptotic events were very low in patients with self-healing

lesions (76). Thus, cell death of some T cells could be involved in the failure of cell mediated immunity, responsible for severe immune-depression in the various forms of leishmaniasis.

The amount of data available suggests that CD8⁺ T cells are involved in leishmanicidal activity, and that this activity is modulated by IFN- γ and IL-2 secretion (77). Many reports have highlighted a variety of effector mechanisms by these CD8⁺ T cells, which involves the direct cytotoxic activity, through the perforin/granzyme pathway, and the induction of apoptosis, via Fas/FasL interaction (76-78). They also secrete various cytokines (especially IFN- γ) and chemokines (especially the ones belonging to the C-C group, i.e. Regulated upon Activation, Normal T-cell Expressed, and presumably Secreted [RANTES]), that might have a key role in acquired cell-mediated immunity against the parasite, i.e. by promoting CD4⁺ T_H1 cell development (78).

Perforin/granzyme-mediated direct cytotoxicity, as well as induction of apoptosis of parasitized macrophages, are candidate mechanisms employed by CD8⁺ T cells in their effort to limit parasite multiplication (76).

Post Kala Azar dermal leishmaniasis

In most cases of PKDL, parasites or parasite antigens were observed in all the lesions and were able to induce the formation of an inflammatory infiltrate, consisting of a mixture of macrophages, lymphocytes and plasma cells (79). In patients who had high IFN- γ responses to *Leishmania* antigen *in vitro*, compact epithelioid granulomas were formed (80). The inflammatory cells were mainly CD3⁺, and IL-10 was the most prominent cytokine found in the lesions (80). However, IFN- γ was found in all lesions in varying amounts, whereas the presence of IL-4 was variable (79-80).

CONCLUSIONS

There is a correlation between the clinical outcome of *Leishmania* infection and the cytokine response profile. While a protective immune response against *Leishmania* has been clearly identified as being under the influence of a type-1 response and IFN- γ production, the precise role of T_H2 cytokines in non-healing infections requires careful exploration (see

the summary in Table I).

IL-4 and IL-13 (T_H2 cytokines) can promote disease progression in appropriate models of CL, whereas the role of IL-4 is not well defined in VL, and also its ability to enhance protective type-1 response has been described. Thus the T_H1/T_H2 paradigm of resistance/susceptibility to intracellular parasites is an oversimplification of a more complicated network of regulatory/counter-regulatory interactions. The role of IL-4 in influencing *Leishmania* infection awaits a re-evaluation according to the *Leishmania* species involved, to the host organism used and to the tissue site examined. Moreover, the presence of antigen specific regulatory T cell subsets may provide an environment that contributes to the balance between T_H1 and T_H2 cells.

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