## MANAGEMENT OF KALA-AZAR IN BIHAR: SCOPE FOR VACCINE DEVELOPMENT

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#### **Abstract**

#### Keywords:

Visceral leishmaniasis, Prophylactic vaccine, Therapeutic vaccine

Leishmania causes a spectrum of diseases globally ranging from self-healing cutaneous to non-healing mucocutaneous and fatal visceral infection forms. Visceral leishmaniasis (VL), caused by the parasite Leishmania donovani, widely affects resource-poor populations in Bihar, India. Post kala-azar dermal leishmaniasis (PKDL) is a sequel of VL. People with PKDL, whose number is gradually on the rise in India, are considered to be a silent parasite pool for transmission of VL. Despite many options, treatment of VL in India is far from satisfactory due to high incidence of PKDL development, drug-resistance, relapse, drug-toxicity, long-dose regimens, need for parenteral administration and high cost. Currently, there is no approved vaccine for human use in leishmaniasis. However, several leishmaniasis vaccine candidates are at various stages of development. Vaccines such as Leish-F1, F2 and F3, are designed based on selected Leishmania antigenic epitopes, have entered the phases of clinical trials. Other groups, including the Sabin Vaccine Institute in collaboration with the National Institutes of Health are working on development of a recombinant Leishmania antigen vaccine with selected sand fly salivary gland antigens targeting to elevate the host immunity status. The state of Bihar in India is heavily affected by VL accounting for over 90% cases of India. In this review, we will discuss the strategies of Leishmaniasis vaccine development, details of vaccines in the developmental pipeline and about future perpectives of an ideal vaccine for effective management of Kala-azar in Bihar.

#### Introduction

The Leishmaniadonovani complex (comprising L. donovani, L. infantum and L. chagasi), an obligate intracellular protozoan parasite, is responsible for a fatal progressive systemic disease- visceral leishmaniasis (VL), commonly referred to as "kala azar" or "black fever". Leishmania parasite follows a digenetic lifestyle by remaining either as extracellular and flagellated promastigotes within sand fly or as intracellular and non-flagellated amastigotes within the mammalian host. The parasites mainly affect mononuclear phagocytic cells of the reticuloendothelial organs, viz. liver, spleen, and bone marrow. However, they are usually disseminated to other visceral organs (gut, lung, etc), as well as the skin, particularly seen in L. infantum-affected areas [1]. Persistent fever, hepatosplenomegaly, pancytopenia, and hypergammaglobulinemia are the characteristic features of human VL [2]. The case fatality rate of this disease is 100% because the patient dies in the absence of treatment. Available chemotherapeutics have several demerits, and their usage is also limited due to the fact that only ailing individuals with clinical symptoms are benefited.

Human VL occurs in a more heterogeneous form with varied chronicity. Only few infected individuals develop clinical symptoms, and on this basis, they are the ones who are subjected to antileishmanial chemotherapy. However, the majority of the endemic population remains with subclinical infection or as asymptomatic carriers (~6–10 times more than the number of VL patients) and their role in disease transmission is yet to be confirmed [3]. Moreover, the immunocompromised state of VL patients makes them susceptible to other secondary infections.

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Emergence of parasite resistance following drug treatment and increased reporting of disease relapse is another major problem in the process of elimination. Furthermore, because the treatment end point is not defined, as patients often fail to report back 6 months posttreatment, it is difficult to ascertain the complete cure from the disease.

Occasionally, approximately 10%–50% of VL patients, during or after the treatment, develop post kala azar dermal leishmaniasis (PKDL), which is characterized by maculopapular or nodular skin rashes [4]. It is commonly seen in East Africa and the Indian subcontinent and requires prolonged and expensive treatment. The role of PKDL in transmission of VL is debatable. [5](Desjeux et al., 2013).

#### **Epidemiology of VL in India**

In India, visceral leishmaniasis (VL) or, Kala-azar is endemic in the four north-eastern states: Bihar, Jahrkhand, Uttar Pradesh and West Bengal. Present spatial spared of VL endemicity limited to 54-districts of these four states. Bihar is epicenter; 33 out of 38 districts are VL affected, of which 10 districts (i.e. Araria, East-Champaran, Gopalganj, Muzaffarpur, Purnia, Saharsa, Siwan, Saran, Sitamrhi and Darbhanga) are highly endemic (report 200 or, more cases annually) and contribute to more than 70% cases of the state [6]. Bihar alone contributes more than 70% of the total Indian cases annually. In 2017, 71.7% of the total Indian cases were reported from the Bihar state only, where as 23.6 % in Jharkhand, 2% in Uttar Pradesh and 2.7% in West Bengal. An approximately 99-million people are at risk in the 1200 villages over the 458 blocks [7]. From 2002 to 2017, a total of 2, 80,780 VL cases with 1,380deaths were reported in Bihar; annul cases ranged from 9505 in 2002 to 4,127 in 2017. Duringthis period,two rising trends were observed (i.e. one in 2002-2007 and another in 2009-2011) [Fig 1]. The highest peak of VL cases occurrence was found in 2007 (i.e. 37, 822). There is a sharp decline in the mortality rate, from 158 in 2002 to 0 in 2017. A decreasing trend was observed in 2012 and 2011 onwards for cases occurrence and mortality, respectively [Fig 1]. Annul case incidences (i.e. 10,000 populations) among the endemic districts varied from 0.01 to 15.44  $(\text{mean}\pm\text{SD} = 1.64 \pm 2.66)$ . The cumulative hot-spot map shows the spatio-temporal distribution of VL-endemicity level throughout the state at four years interval between 2002 and 2017 [Fig 2]. A persistence high level VL cases occurrence was observed in the center and north-eastern part of the state between 2002 and 2013. However, in 2014 to 2017 intensity of VL cases occurrence ranged between low and moderate levels throughout the state and it indicates a good progress in the VL elimination programme of Bihar state to achieve the elimination target in the near future. Although the number of VL incidence have been decreased and the mortality rate has been ensured 'zero' reporting at all the endemic zones of the state, the increasing trend in the PKDL cases occurrence could be a major concern point for an epidemic outbreak/resurgence of the new VL cases in the future [Fig 3] [8.9]. This will not only disrupt the progress in VL endemicity control in the region but also make the situation difficult to sustain the disease occurrence in the low endemic or, non-endemic regions.



Figure 1: Number of VL cases and deaths occurred in Bihar state between 2002 and 2017. Data sources: Bihar State Health Society and the District Malaria Office, Sheikhpur, Patna (Bihar, India).

#### Brief outline of the Immunobiology of VL

The host immunology plays a key role for defining the fate of the disease in VL. Therefore, development of a successful vaccine for VL needs to target the immune system of the host. The major innate immune cells that play a significant role in defense against *Leishmania*are neutrophils, macrophages, and dendritic cells (DCs). When the female sandfly sucks blood meal from the vertebrate hosts, flagellate metacyclic forms of *Leishmania*are delivered along with sandfly salivary ingredients into the skin of the hosts [10]. Initially, the promastigotes are taken up by the neutrophils at the site of infection. Following apoptosis in these infected neutrophils, the released parasites infect neighboring macrophages [11,12]. These macrophages are recruited by the chemotactic properties of the proteophosphoglycans, delivered to the infection site by the vector at the time of its blood meal [13,14]. *Leishmania*parasites are phagocytosed following binding to C3b after which promastigotes get converted to the amastigote form [15]. Infection is established following phagocytosis in macrophages, which is determined by several survival strategies of the parasites, most prominent of which is the modulation and attenuation of immune responses. *Leishmania*parasite suppresses the release of Th1 associated cytokines which in turn restrains DCs to present the parasite-specific antigens to the T cells. Thus preventing the activation of the acquired immunity, the parasite gains survival advantage[16, 17].

The induction of CD4+ Th1 cell responses against parasite antigens is crucial in controlling primary infection, when cytokines such as IFN- $\gamma$  induce nitric oxide production by activated phagocytic cells able to kill internalized parasites. Concomitantly to the role of CD4+ T cells, CD8+ T cells also contribute to protection against disease, and have an important role in controlling primary infections by increasing the Th1 response through a mechanism

dependent on IFN- $\gamma$  production. In contrast, cytokines such as interleukin-4 (IL- 4), interleukin-10 (IL-10), interleukin-13 (IL-13), interleukin-18 (IL-18), and transforming growth factor beta (TGF- $\beta$ ) represent disease promoting molecules which inhibit the Th1 response, contributing to the deactivation of infected macrophages and, consequently, to the development of disease [18]. Although, VL initially was thought to be associated with a Th2-type immune response seen as elevated levels of IL-4 and/or IL-13 [19, 20], some recent studies implicate that there is not a clear Th2 skewing in human VL[21-23]. Typically VL is associated with increased production of multiple and primarily pro-inflammatory, cytokines and chemokines. VL patients have been found to have elevated plasma protein levels of IL-1, IL-6, IL-8, IL-12, IL-15, IFN $\gamma$  inducible protein-10 (IP-10), monokine induced by IFN $\gamma$  (MIG), IFN $\gamma$ , and TNF $\alpha$  [20-22]. An elevated level of IFN $\gamma$  mRNA has been found in the spleen and bone marrow during the acute phase of infection [23]. These observations suggest that development of VL is not driven by Th2 skewing per se, but that other mechanisms also contribute to the pathogenesis of VL [20-23].

Clinical studies strongly implicate the role of IL-10 in many of the immunologic defects associated with kala-azar [23]. Patients with active VL have elevated levels of IL-10 in serums well as enhanced IL-10 mRNA levels in spleen, lymph nodes, and bone marrow [20, 24]. The main disease-promoting activity of IL-10 in VL is probably conditioning host macrophages for enhanced survival and growth of the parasite. IL-10 can render macrophages unresponsive to activation signals and inhibit killing of amastigotes by down-regulating the production of TNF $\alpha$  and NO [20, 25]. In human VL,inhibition of IL-10 enhance the IFN $\gamma$  response by antigen-stimulated PBMC, and neutralization of IL-10 in VL serum inhibit *L. donovani* replication in macrophages [20, 25].



**Figure 2:** Spatio-temporal distribution of cumulative VL cases occurrence at four years interval from 2002 to 2017. Annual VL incidence rate per 10,000 population calculated at district level and the cumulative spatio-temporal distribution map derived through inverse distance weighted (IDW) interpolation technique in ArcGIS software v9.1 (ESRI Inc., Redlands, CA, USA). Propensity of VL endemicity levels (i.e. low, very low, medium, high and very high) visualizes using the intensity of different colors (i.e. yellow and brown) throughout the Bihar state.

Data sources: Bihar State Health Society and the District Malaria Office, Sheikhpur, Patna (Bihar, India).

Although the classical Th1/Th2 paradigm of resistance/susceptibility appears to be valid during CL, a mixed Th1/Th2 response is required for disease control during VL [26]. However, it is yet to establish a clear Th1/Th2 paradigm for curative and preventive response against both CL and VL. Moreover, for ML, the disease manifestation is largely due to inflammatory response than due to parasite burden. Therefore, conventional Th1/Th2 paradigm does not apply to ML. It has been found that Treg (CD4+CD25+ regulatory T cells) as well as Th17 (other subsets of T cells) cells, play a significant role in disease outcome in both CL and VL, their role in ML is much more complicated. Studies with *Leishmania major* and*L. infantum*have shown a protective role of IL-17 as well as IL-22 (Th17 cytokines) against intracellular parasites [27, 28]. Recently it was shown that when recombinant IL-17 or IL-23 was administered to mice it caused a considerable containment of parasite load in infected organs with significant production of factors such as IFN- $\gamma$ , nitric oxide, etc. Thus, this study demonstrated the association of Th17-based cytokines in providing protection against the disease [29].

A lower CD4/CD8 ratio has been described in VL PBMC. In human VL a splenic influx of both CD4 and CD8 T cells are observed implicating that CD8 cells may survive better than CD4 cells in VL patients[20], though in the later stage of the disease leukocytes are scanty and plasma cells and macrophages predominate in the spleen [30]. In mouse models of VL, CD8 cells play important role in controlling *L. donovani*/*L. infantum*infection in liver through their cytolytic activity [31-34]. Moreover CD8 T cell response provide good immune response following ©Indian JMedResPharmSci

vaccination.[35](Stager et al., 2000)As an important constituent of the immune system, Treg cells are known to regulate immune response of other cells. These cells were observed to be present in human cutaneous lesions [36]. Increased expression of lesional FoxP3 and IL-10 during progressive L. major infection in a murine model and similarly during Leishmaniabraziliensisinfection in human patients suggest the disease-promoting role of these regulatory cytokines [23]. The preliminary data suggest that despite Th1 polarization production of IL-10 and Treg cells is associated with delayed healing of CL [37]. Apart from T-cell subsets (Treg and Th17) other than conventional T cells, the role of innate immune response has been essentially linked to disease outcome. In fact, engagement of the macrophage toll-like receptors (TLRs) by the parasites has not only shown to improve phagocytosis but also lead to the killing of the parasites due to triggering of NF-kB transactivation and concomitant production of the downstream mediators including pro-inflammatory cytokines. For instance, TLR9 activation has been found to be beneficial for the host against these parasites. But this situation may not be true for all the TLRs. Lipophosphoglycan, a TLR2 agonist has been shown to have antagonizing effect on TLR9 mediated signal cascade in host macrophage, which in turn facilitates parasite survival [38, 39]. Studies showed that treatment with TLR4 and TLR9 agonists decreased the disease severity following challenge infection with L. major in BALB/c mice [40]. However, in human VL, comparison of mRNA expression levels between pretreatment and posttreatment splenic aspirate samples showed considerably more TLR2 and TLR4 expression but no change in TLR9 expression during L. donovani infection [41]. Furthermore, the later stages of L. donovani infection rendered tolerance to macrophages, leading to incapability for the production of inflammatory cytokines like tumor necrosis factor (TNF)- $\alpha$  and interleukin (IL)-1ß in response to TLR stimulation.[42]. Despite advances, achieving a comprehensive and clear picture of the immunobiology of leishmaniasis is still required to develop effective interventions such as typespecific vaccine and immunotherapy for leishmaniasis.

#### Feasibility of suitable Vaccine against Visceral Leishmaniasis (VL)

The World Health Organization, in 2005, has initiated elimination campaigns in VL-endemic regions through early case detection, followed by complete treatment with a target to reduce the number of diseased individuals in the endemic areas. This drive has successfully saved many lives through the administration of antileishmanial drugs. The only control measure for VL presently available is chemotherapy, which includes pentavalent antimonials (meglumineantimoniate and sodium stibogluconate [SSG]), amphotericin B and its liposomal formulation, paromomycin, and miltefosine either alone or in combination [2]. The chemotherapeutics used against VL have been found to be very effective but with several demerits, such as low efficacy, lesser availability, toxicity, high cost of treatment (drugs and hospitalization) and requirement of longer regimens with the invasive route of administration (parenteral) [43]. The achievement of this goal is difficult with chemotherapy alone due to emergence of drug resistance [44]. Furthermore, the presence of asymptomatic individuals in hyperendemicareas might have resulted in disease transmission, thereby creating hurdles for the current VL control program. Hence, there is an urgent need for alternative treatment strategies to control the kala-azar. Over the past few decades, researchers have been trying to develop a vaccine as a cost-effective treatment strategy against VL [45]. It was evident that patients recovered from Kala-azar develop strong immunity, making them resistant to subsequent clinical infections, which indicates that prevention of Kala-azar through either prophylactic or therapeutic vaccination is feasible [46]. Identification of appropriate vaccine candidates through a proper understanding of the immunobiology and pathogenesis of the disease is required for the success of the vaccine development program [41]. Additionally, selection of either a suitable adjuvant or immunomodulatory or an effective delivery system can further boost the immune responses and generate long-lasting immunity [47].

#### **Prophylactic Antileishmanial vaccine**

Although initial studies suffered with objectionable side effects resulting into clinical complications, later various leishmanial antigens have been explored as promising vaccine candidate for leishmaniasis. These antigens included killed or live attenuated *Leishmania* parasite (first generation), recombinant Leishmania proteins (second generation), DNA encoding Leishmania proteins (third generation), and immunomodulators[48-51]. Admittedly, at present there is no vaccine available for human use against leishmaniasis.Many scientists are investigating the possibility of vaccination against leishmaniasis[49, 52]. The major challenges in the development of vaccine against leishmaniasis are the complexity associated with the antigenicity, uneven response by the host, variability in the

different species of the Leishmania, and cost associated with the development [51]. The approaches for prophylactic vaccine studies are summarized in the Table 1.



**Figure 3:** Number of PKDL cases occurred in Bihar state between 2012 and 2017. A 5.2 folds increase observed in occurrence of PKDL cases from 2012 to 2017. **Data sources:** Bihar State Health Society and the District Malaria Office, Sheikhpur, Patna (Bihar, India).

#### **Options for Therapeutic Vaccines against VL**

Therapeutic vaccines are thought to be essential in case of persistent chronic infections, because in such cases, either intracellular parasites evade the host immune system establishing themselves in a more secured way or the available control intervention is ineffective. In case of VL, the majority of inhabitants in disease affected areas either serves as endemic healthy individual or asymptomatic individual, who serve as the reservoir of parasites or may become symptomatic in future. Hence there is a need for a therapeutic vaccines that can be effectively used to stimulate patients own immune defense system in this endemic population, thereby controlling the progression of the disease. The therapeutic strategy involves the use of biological molecules (whole or their components) in combination with either adjuvants or drugs to modulate the immune responses of *Leishmania*-infected individuals toward the protective type. Therefore, this strategy, which restores or induces an effective immune response without any side effects, could be a promising alternative to conventional chemotherapeutics. The various therapeutic approaches of vaccination against VL according to three generation has been summarized in the following Table 2.

#### **Challenges in vaccine development**

The potential approach for the effective control and complete eradication of any infectious disease is vaccination. Over the past two decades, immunotherapy, either alone or in combination with chemotherapy, has been developed

as an additional approach to combat leishmaniasis. Lifetime immunity against reinfection manifests possibility of developing an effective vaccine (prophylactic and therapeutic) against leishmaniasis. Vaccine development project for leishmaniasis seems to attract less industry people causing a serious crunch of funds. According to the G-Finder, over US\$ 66 million has been granted for research and development of vaccine, preventative, and therapeutic, against leishmaniasis largely from chief public sector and charitable trusts (from the year 2007 to 2013). Some of the major funding sources are Carlos Slim Foundation, Bill & Melinda Gates Foundation, Wellcome Trust, Indian Council of Medical Research, European Commission, Institute Pasteur, German Federal Ministry of Education and Research (BMBF), and the U.S. National Institutes of Health (Policy Cures. G-Finder; 2015)[104]. Leishmaniainfected individuals gain considerable lifelong immunity to reinfection, suggesting the feasibility of vaccination. However, regardless of many potential vaccine candidates, translation of these to develop a human administrable antileishmaniasis vaccine is still arduous. There are never ending debates regarding the choice of suitable antigens as vaccine candidate with lower toxicity and greater efficacy and immunogenicity despite of studying a series of antigens as described in Table 1 and 2. Leishmania infection follows a complex clinical outcome varying from the cutaneous to visceral form as the parasite is equipped for generating an extensive assortment of atypical and uncommon variations. The virulence factors as well as in the immune responses induced by the different strains and species of Leishmania is not fully known. A better understanding of the immunobiology and vaccine (prophylactic and therapeutic) development prerequisites for the different forms of leishmaniasis will provide tools that can be exploited to overcome the virulence dynamics of Leishmania species. Preliminary studies are mainly done in mouse and hamster model hence the immune responses leading to protection in humans differs due to lack in correlation to the immune response in the animal model.

#### The fore-runners in vaccine development

Chemotherapapeutic options are the main treatment for all three major forms of leishmaniasis. However, several factors such as high costs, toxicity, and long-term complicated regimens are hurdles for the chemotherapeutic options. The elimination of leishmaniasis also depends on sand fly vector control approach through indoor residual spraying [IRS] in many areas. Emergence DDT resistancein highly endemic areas has jeopardized IRS leading to large growth of sand flies and potential parasite transmission. Therefore, the development and delivery of an ideal long-term immunity providing vaccine may represent as the most cost effective means of controlling and/or eliminating VL in India.

Ideally, an effective vaccine against leishmaniasis should provide long-lasting immunity and protect broadly against both VL andCL. Many leishmanial antigens have been researched extensively for their use as candidate vaccines in preventative and therapeutic indications; either by delivery as DNA or directly as recombinant proteins. Many such vaccine candidates are at various developmental stages of preclinical testing. The first vaccine candidate that made into phase I and II clinical trials is known as LEISH-F1, developed at The Infectious DiseaseResearch Institute (IDRI, Seattle, WA) [105]. LEISH-F1, is a fusion of three tandem polypeptides in monophosphoryl lipid A-stable emulsion (MPL-SE). Phase I trials showed satisfactory safety and immunogenicity profile in endemic and nonendemic populations of the United States, Colombia, Brazil,Peru and India. Interestingly, LEISH-F1 demonstrated shortened time to cure when used with chemotherapy [106, 107]. After the encouraging results of LEISH-F1, IDRIdesigned LEISH-F2 without a histidine tag at the N-terminus, and progressed it into the phase I and a phase II clinical trials. Further changes were made in LEISH-F2 by fusing, in tandem, the open reading frames of nucleoside hydrolase (NH) from *L. donovani* and sterol 24-c-methyltransferase (SMT) from *L. infantum*, with a TL4 based adjuvantglucopyranosyl lipid A to form LEISH-F3. This trial results were promising that found LEISH-F3 to be safe and immunogenic at lower doses.

Other mention-worthy group for vaccine development is the Sabin Vaccine Institute ProductDevelopment Partnership (Sabin PDP). They are exploring prototypecombination vaccines comprised of recombinant proteins encoding sand fly salivary gland antigens and *L. donovani* NH36expressed in either yeast or bacteria. Although there is some encouragingdata from the animal models for this vaccine, the Sabin PDP is still in the R&D phase. Besides this, some other groups of investigators, including the European Multivalent Vaccine for HumanVisceral Leishmaniasis (MuLeVaClin), are working with recombinant protein-based, DNA-based, and heterologous primeboost vaccine strategies for VL in pre-clinical models [108, 109].

The epidemiological scenario of VL makes its elimination a realistic goal. Lastly, in the vaccine development scenario, it is very critical to develop an effective vaccine delivery system and adjuvant in order to achieve a successful *Leishmania* vaccine. Besides vaccine development, we should also focus into repurposing of approved drugs and nutraceuticals to be a great new option for treatment of VL. Moreover, increased financing support from major funders will be a critical need to advance candidate vaccine studies for human use in the target population in India.

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#### Table 1: Prophylactic vaccine used against Leishmaniasis Antigen Vaccine Animal model Adjuvant refer approaches 1<sup>st</sup> generation anti-leishmanial vaccine Merthiolated sound-disrupted BCG Killed vaccine Dogs [53, 54] Leishmaniabraziliensis Killed L. braziliensis Killed vaccine Dogs Saponin [55, 56] KBMA Leishmaniainfantumchagasi Killed vaccine Mouse [57] p27 gene knockout L. donovani parasites Live [58] mouse \_ attenuated Live mutants of Leishmania lacking genes like [59] Live mouse dihydrofolate reductase, biopterin reductase, attenuated and cysteine proteases (CPs) Ascorbic acid-deleted live mutants of L. Live Mouse \_ [60] donovani attenuated Centrin-deficient parasites of L. donovani Live [61] Dogs \_ attenuated

**Prophylactic Vaccine against VL** 

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2<sup>nd</sup> generation anti-leishmanial vaccine

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	I	1	1	1
Chimeric peptides containing HLA-A2	Peptide			[62]
restricted epitopes from six immunogenic L.	Vaccine			
major proteins(CPB, CPC, LmsTI1, TSA, LeIF				
and LPG-3)				
L. donovanisurface GP63	Recombinant	Mouse, Human	MPL-TDM, CGP-	[63, 64]
	protein	,	ODN	L, - ]
	vaccine		ODIT	
Soluble leichmonial antigons of L donovani	Fractionated	Mouso	MDI TDM	[65]
soluble leisinnannar antigens of L. ubnovani	Vaccina	Wibuse		[05]
		TT / 1		[(()]
L. donovanip45 (rLdp45)	Recombinant	Hamster and	-	[66]
	protein	Human		
115 kDa soluble serine protease	Fractionated	Mouse	IL-12	[67]
	vaccine			
L. braziliensispromastigote proteins	Fractionated	Dog	Saponin	[68]
	vaccine			
	Recombinant	Mouse	CpG-ODN	[69]
Leishmania maior ribosomal protein L3 or L5	protein		1	
Leishmune (purified L donovani fraction FML)	Fractionated	Dog	Saponin	[70]
Leisinnune (purneu E. uonovani maction i ME)	Vaccine	Dog	Supolini	[/0]
Leich Tee (L. deneugni emerticate energific	Decembinant	Dec	Cononin	[71]
Leish-Tec (L. aonovani amastigote-specific	Recombinant	Dog	Saponin	[/1]
protein A2)	protein	TT .		(70)
Cocktail of L. donovani CPs types I, II, and III	Recombinant	Hamster	MPL-TDM	[72]
	protein			
	cocktail			
	vaccine			
Recombinant L. tarentolae secreting PpSP15	Recombinant	Mouse	CPG-ODN	[73]
	vaccine			
L. infantumacidic ribosomal P0	Recombinant	Hamster	-	[74]
	protein			
Cocktail of rCDV-LACK rCDV-TSA and	Recombinant	Dog	-	[75]
rCDV-I mSTI1	protein	205		[,5]
	cocktail			
	Vegging			
	Vaccine	Turner	Dal (last's sa	[7](]
	Peptide	I ransgenic	Poly (lactic-co-	[/6]
Chimeric peptides containing HLA-restricted	vaccine	mice	glycolic acid	
epitopes from three immunogenic <i>L. infantum</i>			nanoparticles	
proteins (CPA, histone H1 and KMP11)			and/or MPL-A	
3 <sup>rd</sup> generation anti-leishmanial vaccine				
L. donovanisurface GP63	DNA vaccine	Mouse, Human	MPL-TDM, CGP-	[63, 77]
	and T-cell		ODN	
	epitope DNA			
	vaccine			
Cocktail of L major CPs type I II and III	Cocktail DNA	Mouse	MPL-TDM	[78]
Coektun of L. major Cr 5 type 1, 11, and 111	vaccine	1100.50		[,0]
Lastopasillus lastisovpressing LACV and	Pacombinant	Mouso		[70]
Laciobacillus lacusexpressing LACK and	Recombinant	wouse		[/9]
mouse IL-12	vaccine			[00]
Recombinant L. donovani protein disulfide	DNA vaccine	Hamster and	-	[80]
isomerase		Human		1

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Leishmaniatarentolaeexpressing L. donovaniA2	Recombinant	Mouse, Dogs	-	[81, 82, 83]
antigen along with CPs [CPA and CPB without	vaccine and			
its unusual C-terminal extension (CPB-CTE)]	DNA vaccine			
L. major TSA	DNA vaccine	mouse	Aluminium	[84]
			phosphate	
Leishmaniaamazonensisiron superoxide	DNA vaccine	mouse	-	[85]
dismutase				
T-cell epitope of KMP11_CPA_CPB_EE1a	Multiantigenic	Mouse	-	[86]
and TSA (I FISHDNAVAX)	T-cell enitone	Wiouse		[00]
	fusion $DNA$			
	Iusion DINA			
1	vaccine			

Table 2						
Antigen	Vaccine approaches	Animal model	Adjuvent	References		
1 <sup>st</sup> generation						
L. donovani/ amastigote UR6	Live/sonicated parasite	Hamster	-	[87]		
L. donovani/ amastigote UR6	Live/sonicated parasite	BALB/c mice	-	[88]		
Alum/Autoclaved L. major	killed parasite	Human PKDL patients (field trial in Sudan)	Bovine Calmette- Guerin	[89]		
L. donovani promastigote	Attenuated parasite	BALB/c mice	-	[90, 91]		
2 <sup>nd</sup> generation						
Fucose mannose ligand (FML)	glycoprotein	BALB/c mice	saponin	[92]		
FML	glycoprotein	Mongrel dogs (experimental; Brazil)	saponin	[93]		
Leishmune <sup>®</sup> (FML)	glycoprotein	Mongrel dogs (experimental; Brazil)	saponin	[94]		
L. infantum-derived Fraction-2	Fractionated protein vaccine	Naturally infected dogs (France)	-	[95]		
NH-DNA	glycoprotein	BALB/c mice	Aqueous garlic extract	[96]		
Leish-110f	Polyprotein vaccine	Naturally infected mongrel dogs (Brazil)	Monophosphoryl Lipid A plus squalene (MPL- SE)	[97]		
Complete Soluble Antigen (CSA)	Protein vaccine	BALB/c mice	-	[98]		
Soluble Leishmania Antigen	Protein vaccine	BALB/c mice	Pulsed Dendritic	[98]		

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(SLA)			Cell		
Leish-111f	Polyprotein	Naturally	MPL-SE	[99]	
	vaccine	infected dogs			
		(Brazil)			
Leishmune	Protein vaccine	Naturally	-	[100]	
		infected dogs			
		(Brazil)			
recombinant cysteine proteinase	recombinant	Naturally	-	[101]	
from <i>L</i> . ( <i>L</i> .)	protein vaccine	infected dogs			
infantumChagasirLdccys1+ P.		(Brazil)			
acnes					
GRP78	Protein vaccine	BALB/c mice	Monophosphoryl	[102]	
			Lipid A (MPLA)		
3 <sup>rd</sup> generation					
LEISHDNAVAX	DNA vaccine	C57BL/6J mice		[102]	