Chapter from the book *Advances in the Study of Genetic Disorders*
Downloaded from: http://www.intechopen.com/books/advances-in-the-study-of-genetic-disorders

Interested in publishing with InTechOpen?
Contact us at book.department@intechopen.com
1. Introduction

The leishmaniasis is a group of diseases caused by protozoan haemoflagelates of the genus *Leishmania*\(^1,2\). These parasites belong to the family of the *Trypanosomatidae* (order *Kinetoplastida*) and are closely related to the Trypanosomes\(^3\). Despite enormous efforts, it has proved difficult to predict the exact scale of the impact of leishmaniasis on public health, since many cases remain unreported or misdiagnosed\(^4\). It is estimated that approximately 12 million people are currently infected and a further 367 million are at risk of acquiring leishmaniasis in 88 countries, 72 of which are developing countries and 13 of them are among the least developed in the world\(^1,\ 4\). Hence we can link leishmaniasis to poverty, economic development and various environmental changes such as deforestation, urbanization, migration of people into endemic areas and building of dams etc\(^5\). The annual incidence rate is estimated to be 1 to 1.5 million cases of cutaneous leishmaniasis (CL) and 5,00,000 cases of visceral leishmaniasis (VL); these are the two major clinical types of leishmaniasis\(^6\). The only proven vector of the *Leishmania* parasite is the blood-sucking female sandfly\(^1,7\) of the genus *Phlebotomus* in the old world and *Lutzomyia* in the new world\(^8\).

The insects are 2-3 mm long (one-third the size of typical mosquitoes) and are found throughout the tropical and temperate parts of the world. The sandfly larvae require organic matter, heat and humidity for development and so are commonly found in house-hold rubbish, burrows of old trees and in cracks in house walls\(^9\). The sand flies usually feed at night while the host is asleep\(^10\). There are five most important *Leishmania* species namely *L. tropica*, *L. major*, *L. donovani*, *L. braziliensis brasiliensis*, *L. b. peruviensis* and *L. mexicana* which cause the three forms of the disease dermal CL (oriental sore), VL and mucocutaneous leishmaniasis (Chiclero's diseases and Espundia)\(^11,\ 12,\ 13\). *Leishmania* exhibits a dimorphic life cycle\(^14\) involving two life-cycle stages, the elongated promastigote with free flagellum present in the insect and the intracellular amastigote form\(^15\).
VL, commonly known as kala azar, is characterized by irregular bouts of fever, substantial weight loss, swelling of the spleen and liver, and anaemia (occasionally serious)\textsuperscript{16}. VL was first described in 1903, by Dr. William Boog Leishman\textsuperscript{17, 18, 19}, an English military surgeon and Dr. Charles Donovan\textsuperscript{17, 20}, an Irish physician by identifying the parasites in spleen smears of a patient died of “Dum-Dum fever” i.e. a low degree of fever with hepatosplenomegaly and severe progressive cachexia (wasting), swollen lymph glands, leucopenia, thrombocytopenia with relative monocytosis and loss of hair\textsuperscript{18, 20}. These physicians reported the existence of the parasite for visceral leishmaniasis and so the causative agent acquired the name as \textit{Leishmania donovani}\textsuperscript{21}. This is not to say that leishmaniasis did not exist before 1903, on the contrary. Archibadi in 1922 described an epidemic of Kala-azar, which occurred in the Garo hills of Assam\textsuperscript{22} and in Saudi Arabia\textsuperscript{23, 24} as far back as 1870. Cunningham recorded a similar disease that occurred in 1885, caused by a parasite\textsuperscript{25}, which was later named \textit{Leishmania tropica}, the causative agent of CL\textsuperscript{26}. Nicolle in 1908 reported that mammals including dogs could act as reservoir hosts for the \textit{Leishmania} parasite\textsuperscript{27}. Swaminath \textit{et al} in 1942, proved using human volunteers that the \textit{Leishmania} parasite could be transmitted by the phlebotomus sandflies\textsuperscript{28}. If left untreated, the fatality rate of VL in developing countries can be as high as 100% within 2 years. The recommended drugs for VL are the pentavalent antimonials sodium stibogluconate (Pentostam) and Meglumine Antimoniate (Glucantime)\textsuperscript{29}. Both drugs have been used for over 50 years, and they require long courses of parenteral administration. The treatment has traditionally been unsatisfactory because of drug toxicities, poor responses, multiple disease syndromes and other factors including the recent emergence of antimony-resistant strains\textsuperscript{30}. Side effects of sodium stibogluconate (SAG) include changes in liver function, biochemical pancreatitis, electrocardiogram (EKG) changes, musculoskeletal symptoms, thrombocytopenia etc. Alternative treatments for VL include the polyene antibiotic amphotericin B\textsuperscript{31} that constitute highly effective, less toxic lipid formulations\textsuperscript{32, 33}. In regions of India where there is a high frequency of resistance to antimony, amphotericin B in a dose of 15-20 mg/kg, body wt is administered intravenously (IV) over a period of 30-40 days\textsuperscript{34}. Recently several compounds of herbal origin have been reported to have potency against VL\textsuperscript{35, 36, 37}. Post Kala azar dermal leishmaniasis (PKDL) occurs in India and mainly in Sudan and Kenya in Africa\textsuperscript{38}. Reports of PKDL in China, Iraq and Nepal have also been documented \textsuperscript{39, 40}. In the new world PKDL is extremely rare\textsuperscript{41}. Usually PKDL follows recovery from a Kala azar infection, though less commonly, it has been known to occur in patients who have not suffered previously from Kala azar\textsuperscript{42}. Both the Indian and the African PKDL display similar symptoms. The disease begins with small measles-like lesions (hypopigmented macules, papules or nodules) appearing on the face, and gradually increase in size. Eventually the lesions spread to the upper trunk, arms, forearms, thighs, legs, abdomen, the neck and the back. The multiple lesions can coalesce to form larger lesions and can lead to the gross enlargement of facial features such as the nose and lips, giving an appearance similar to leprosy. The disease is particularly severe if the lesions spread to the mucosal surfaces of the nasal septum, hard and soft palate, oropharynx, larynx or the eye lids and the cornea leading to blindness\textsuperscript{43, 44}. Potentially the lesions can appear on any part of the body. There have been reports of lesions occurring on the glands of the penis and genital mucosa hence the possibility of PKDL being transmitted sexually. In addition to the disease being confused with leprosy, PKDL can also resemble cutaneous leishmaniasis, secondary syphilis and sarcoidosis\textsuperscript{44}. The lesions are usually self-limiting, however those that do not
heal spontaneously within six months have to be treated\textsuperscript{45}. Pentavalent antimonials remain the drugs of choice for treating PKDL. SAG at a dose of 20 mg/Kg of body weight administered intramuscularly for 4-5 months is recommended. In addition Ketoconazole and allopurinol can be given orally to improve response. In antimony resistant cases amphotericin B is an effective replacement\textsuperscript{46, 47}. The reason for incidence of PKDL is still not clear and the evidence is yet to be explored. Several studies indicate that PKDL may develop as a result of genetic disorder during the parasitic disease kala azar. Indian PKDL appears anything between 1-7 years after apparent cure of Kala azar, although longer periods of upto 20-30 years have been reported\textsuperscript{47}. The African form of the disease usually appears within a few months after cure, in most cases within 6 months, on average within 56 days. However it can develop during the treatment Kala azar, in which case the term Para Kala azar dermal leishmaniasis would seem more fitting\textsuperscript{48}. The choice available drug is limited and inadequate\textsuperscript{49}.

Although the geographical distribution of \textit{Leishmania} infection is restricted to the area of distribution of \textit{Phlebotomus} sandflies, Human immunodeficiency virus (HIV) infection may modify the traditional anthroponotic pattern of VL transmission. Very rarely \textit{Leishmania} transmission has been described by alternative means that are also shared by HIV transmission, including blood transfusion\textsuperscript{50} congenital transmissions\textsuperscript{51, 52, 53} and laboratory acquired\textsuperscript{54}. Worldwide, VL mainly occurs in HIV negative individuals more so in paediatric patients\textsuperscript{55}. The association of \textit{Leishmania} with HIV\textsuperscript{56} has lead to a significant shift in the age of people at risk\textsuperscript{57, 58, 59}.

In South West Europe, 75 per cent of HIV sero negative and 80-83 per cent of HIV positive patients seen with VL were men\textsuperscript{59, 60}. \textit{Leishmania}/HIV co-infection is emerging as a serious new disease pattern and is becoming increasingly frequent. Of the 1700 cases of co-infection reported to the World Health Organization from 33 countries worldwide up to 1998, 1440 cases were from South-West Europe: Spain (835), Italy (229), France (259) and Portugal (117). Of 965 cases retrospectively analyzed, 83.2 per cent were males, 85.7 per cent were young adults (20-40 yr) and 71.1 per cent were intravenous drug users\textsuperscript{61}.

Most co-infections in the Americas are reported from Brazil where the incidence of HIV has risen from 0.8 cases per 100,000 inhabitants in 1986 to 10.5 cases per 100,000 inhabitants in 1997\textsuperscript{62}. In India, the HIV/\textit{Leishmania} co-infection has not been extensively studied. The risk of visceralization for HIV positive person, infected with \textit{Leishmania} species typically associated with cutaneous disease, is not much of a problem, since VL is several thousand times more common than CL in India. However, this may be a very serious issue in the Mediterranean countries where CL is very common\textsuperscript{63}.

A majority of HIV/\textit{Leishmania} co-infected cases show classical features of VL. These co-infected patients may also have other features, viz., atypical location due to decreased cell mediated immunity (CMI) \textsuperscript{64, 65}, parasitic dissemination to skin, cutaneous and reticuloendothelial system (RES), a chronic and a relapsing course, poor drug response and lack of anti-leishmanial antibodies.

The incubation period is variable and may be age related\textsuperscript{66, 67}. Other concomitant opportunistic infections are diagnosed in 42-68 per cent of HIV-positive patients\textsuperscript{68}. Fever, pancytopenia, hepatosplenomegaly are common. Classically, splenomegaly may be less in HIV positive patients\textsuperscript{69}. Constitutional symptoms (asthenia, anorexia, weight loss etc.) are seen in 50-70 per cent of patients and lymphadenopathy in 15-60 per cent of patients. VL with HIV infection may present as pyrexia of unknown origin (PUO). Other opportunistic
infections like mycobacterial infection, cytomegalovirus (CMV), pneumonias and Acquired
immune deficiency syndrome (AIDS) related neoplasms may also occur\textsuperscript{70, 71}.
Gastrointestinal (GI) symptoms are among the most frequent complaints\textsuperscript{72, 73, 74}. Leishman-
Donovan (LD) bodies have been identified in up to 50 per cent of such patients. The commonest site of involvement is the jejunum\textsuperscript{75, 76}. Endoscopy and routine biopsy are important tools in the diagnosis. The symptoms may include diarrhoea, malabsorption, and hypoalbuminaemia and weight loss. There may be erosive gastro-duodenitis, ulcers and colony lesions.

Cutaneous involvement\textsuperscript{77} may appear in the skin with Karposi sarcoma, Herpes simplex or
Zoster. \textit{Leishmania} may be associated with dermatofibroma, psoriasis, Reiter’s syndrome,
bacillary angiomatosis, cryptococcosis and oral aphthous ulceration. It may also present as
dermatomyositis like eruption\textsuperscript{78, 79}.

Respiratory tract involvement occurs in alveoli and pulmonary septa in 75 per cent of
patients with VL\textsuperscript{80}. They could present with pulmonary tuberculosis and pneumonia, more
commonly Pneumocystis carinii pneumonia (PCP). The symptoms could be cough, breathlessness,
haemoptysis and excessive sputum production. Renal involvement can occur. Glomerulonephritis with mild proteinuria, haematuria and even acute renal failure have been reported. Tubulointerstitial damage can also occur\textsuperscript{81}.

Central nervous system (CNS) involvement is very common in the late stages. Pandey et al\textsuperscript{82}
reported cases in which HIV-\textit{Leishmania} co-infection was associated with pulmonary
tuberculosis and tuberculoma in the brain neurocysticercosis and tuberculous meningitis.
AIDS dementia complex occurs in the late stages and may lead to early death. In such cases,
colour of differentiation 4 (CD4) count has been reported to be as low as &lt;50 cells/mm\textsuperscript{3}.
Pancreatic, pulmonary, pleural, laryngeal, adrenal, pericardial, myocardial and lingual
leishmaniasis have also been reported. Mucocutaneous leishmaniasis appears in 2-3 per cent
of VL-HIV co-infected patients\textsuperscript{83-89}.

Diagnosis is quite difficult as only 40-50 per cent of VL/HIV co-infected cases have a
positive \textit{Leishmania} serology\textsuperscript{90}. This percentage is inversely proportional to CD4 cell
depletion. Anti-\textit{Leishmania} antibodies in HIV-positive patients are 50 times less than those in
HIV-negative patients\textsuperscript{91}. Therefore, there may be many false negative tests. The direct
examination of amastigotes in the splenic and bone marrow aspiration has been the gold
standard. Detection of \textit{Leishmania} antigens by Western blot in the urine samples is being
tried. Polymerase chain reaction (PCR) techniques requiring blood and tissue samples are
very time consuming\textsuperscript{92, 93}. However, when used in combination with Enzyme-linked
immunosorbent assay (ELISA) and Direct Agglutination Test (DAT) the results are very
couraging. Nested PCR assay has a sensitivity of 95 per cent in the peripheral blood and
100 per cent in bone marrow/ splenic aspirates\textsuperscript{94, 95}.

Certain issues are significant in the management of HIV/\textit{Leishmania} co-infected patients.
Firstly, optimal duration of treatment is to be given. Secondly, the dose has to be monitored
and thirdly, there is frequent relapse.

Sodium antimony gluconate (SAG) has developed resistance and low cure rates of 30-50 per
cent have been reported\textsuperscript{92}. In such patients, the treatment with SAG has to be given for a
longer period\textsuperscript{96}. However, the longer duration of therapy may lead to cardiotoxicity. With
SAG, there are frequent relapses as seen often in Bihar, India\textsuperscript{97}. Amphotercin B has also
been tried and response rate of 60 per cent has been observed but 25-60 per cent of the
patients treated with amphotercin B, are likely to have relapse during the first year after
completion of treatment\textsuperscript{98}. HIV infected individuals are more likely to suffer from drug
related adverse events. In almost all the patients, depending on the CD4 counts (<200/ml) Highly Active Antiretroviral Therapy (HAART) can be given. Amphotericin B can be given at a dose of 1 mg/kg for 15 days. Although lipid formulations are less toxic, they are very costly. Pentamidine is usually not effective and should not be used due to its toxic effects. Oral miltefosine is a promising alternative at a dose of 2.5 mg/kg for 28 days and has been tried by Thakur et al. in six co-infected patients with good results. Besides treatment of VL and administration of HAART, other secondary infections like tuberculosis of the chest, oral candidiasis, CMV infections, Pneumocystis carinii pneumonia (PCP), toxoplasmosis, Karposi sarcoma also need to be treated. Peroxisomes were first detected morphologically; their biochemical functions were characterized sometimes later. Johannes Rhodin, in 1954, described small organelles (0.5 μm in diameter) in mouse kidney cells and identified them as microbodies. Later in 1965, Christian de Duve proposed the name peroxisomes (as they are peroxide-metabolizing organelles), for the distinct organelles isolated by density-gradient centrifugation. These organelles contained enzymes for both the production and disposal of hydrogen peroxide (H₂O₂). H₂O₂ generates reactive oxygen species, which are fatal in all respects. They interact with a variety of cellular macromolecules and lead to variety adverse effects like, membrane damage (lipid peroxidation), DNA damage (strand breakage), alteration in the protein structure (thiol oxidation) and the ultimate consequence may be the loss of cell viability (cytotoxicity and cell death). Catalase is the marker enzyme for peroxisome as it is the most abundant enzyme within the structure. Active catalase is a heme- containing protein with four identical subunits (tetramer). In mammalian liver and kidney cells, peroxisomes are roughly spherical and relatively large (0.5 to 1.5 μm in diameter). In other tissues, they are smaller (0.1 μm) and known as micro-peroxisomes. In mammalian cell, the average number of peroxisomes is approximately 400 that occupy 1% of the total volume of the cell. However, their size, number, and enzyme profile vary between different tissues. In most cases, peroxisomes contain dense or granular matrix containing catalase (About 40% of the total peroxisomal protein is catalase) and a crystalline core composed of enzyme urate oxidase. Peroxisomes have a half-life of about 36 hrs and they divide by fission. The process of formation of peroxisome involves multiple pathways. Lipids must be recruited to form the membrane. Lipid composition of the peroxisomal membrane differs from that of the total cellular profile. Peroxisomal proteins (not glycosylated) are synthesized in the cytosolic free polyribosomes and transported to pre-existing peroxisomes. These include both the peroxisomal membrane proteins (PMP) and peroxisomal enzymes. The multifunctional structures begin through orchestrated reactions of some proteins, called peroxins. These are the critical processes as their defects leave the cells either devoid of peroxisomes or with organelles rendered unable to carry out numerous biochemical and metabolic functions attributed to them; such failings often cause a disease. Approximately fifty different biochemical processes occur exclusively within a peroxisome. Some of the reactions are anabolic i.e. constructive, resulting in the synthesis of essential biochemical compounds including bile acids, cholesterol, plasmalogens (phospholipid analogues), and docosahexanoic acid (DHA), which is a long chain fatty acid that is
Advances in the Study of Genetic Disorders

Component of complex lipids, including the membranes of central nervous system. Other reactions are catabolic i.e. destructive and lead to the lysis of some fatty acids, via \( \beta \)-oxidation including very long chain fatty acids\(^{117-119} \).

Peroxisomes contain several oxidative enzymes, which generate \( \text{H}_2\text{O}_2 \) as a by-product of their reactions. This highly poisonous reactive oxygen species (ROS) is rapidly converted to \( \text{H}_2\text{O} \) through the action of peroxisomal catalase, at least under most circumstances. A number of degenerative diseases are linked to ROS induced alternation in cellular functions\(^{120} \). The reactions are catalyzed by appropriate FAD-linked enzyme: a) urate oxidase, b) xanthine oxidase, and c) L- and D- amino acid oxidases. Flavin oxidases reduce oxygen to \( \text{H}_2\text{O}_2 \) which is then decomposed by catalase\(^{121} \). However the peroxisomal pathway differs from that of the mitochondria in several important respects\(^{122} \).

\( \beta \)-oxidation pathway similar to that of mitochondria is present in peroxisomes and the process is carried out by two distinct groups of enzymes. The classical first group utilizes straight chain saturated fatty acyl-CoA as substrates, whereas the second group acts on the branched chain acyl-CoA\(^{119, 124} \).

Severe effects of peroxisomal dysfunction, which secondarily leads to several human and animal diseases, emphasize the importance of peroxisomes for survival. Depending on the specific defect, clinical manifestations range from the mild to the fatal.

There are about 25 peroxisomal disorders known, although the number of diseases that are considered to be separate, distinct peroxisomal disorder varies among the researchers and health practitioners\(^{125, 126} \). Peroxisomal disorders are subdivided into two major categories. The first category is the disorders resulting from a defect in a single peroxisomal enzyme\(^{127} \). These disorders include hyperoxaluria type I (alanine: glyoxylate aminotransferase), Refsum’s disease (phytanoyl-CoA hydroxylase), X-linked adreno-leukodystrophy (ALDP), rhizomelic condrodysplasia punctata (RCDP) types II and III (dihydroxyacetone phosphate acyl transferase), and the \( \beta \)-oxidation disorders (acyl CoA-oxidase, bifunctional protein, and thiolase). These disorders result from a deficiency in only a single enzyme of the peroxisome, and therefore generally only affect a single peroxisomal metabolic pathway.

Conversely, there is also a set of disorders which results from a malfunction to form intact, normal peroxisomes, resulting in multiple metabolic abnormalities, which are referred to as peroxisome biogenesis disorders (PBD) or as generalized peroxisomal disorders\(^{128} \) and include the Zellweger syndrome (ZS), Neonatal adrenoleukodystrophy (NALD), Infantile Refsum disease (IRD). Rhizomelic condrodysplasia punctata (RCPD) type I.

Several proteins have been identified which play a role in the degradation of peroxisomes\(^{129-133} \). There are a number of reports in the literature describing the histology of the liver, spleen, and lymphoid organs of VL, during human disease and experimental infection\(^{134-136} \). In an experimental model of VL in golden hamsters, histopathological analysis showed\(^{137} \) dissemination of the parasite mainly to liver & spleen. The former organ showed hypertrophy and hyperplasia of Kupffer cells with focal areas of inflammatory infiltration in nodular pattern. Gross examination of the liver indicated hepatomegaly. The spleen disclosed intense proliferation and enlargement of mononuclear phagocytic cells, revealing nodular configuration.

Both biochemical and morphological changes that take place in host peroxisomes during Leishmania infection, was detected\(^{138, 139} \) pointing to the occurrence of a peroxisomal disorder during this parasitic disease. Liver peroxisomes were found to be functionally defective when purified after L. donovani infection. The activities of key enzymes catalase, urate oxidase, dihydroxy acetyl phosphate acyl transferase (DHAPAT) and superoxide dismutase (SOD)
were either deficient or could not be detected after parasite infection. H₂O₂ producing peroxisomal β-oxidation was significantly elevated after 90 days of infection, with concomitant induction of superoxide radical production. Proteolytic activity in infected liver peroxisome was found to be inhibited, pointing to possible uneven processing of peroxisomal proteins. The morphology of peroxisomes after Leishmania infection was impaired. The evidence obtained for Leishmania-induced peroxisomal dysfunction may provide clues to develop new drugs against this parasite, capable of protecting normal function of this ubiquitous host organelle for successful treatment. Calcium is known to be stored in mitochondria and endoplasmic reticulum and is mobilized by second messengers. As one of the vital organelles of eukaryotic system, role of peroxisomes in mobilizing calcium has recently been explored in which peroxisome was identified as a calcium-containing intracellular organelle for its possible candidacy as one of the mediators towards cell signaling.

Immense interest has recently been developed to study peroxisomal properties in more details due to involvement of this mammalian microbody in normal cellular function. It has already been reported that infection due to Leishmania pathogen leads to host peroxosomal damage. As peroxisome is known to be involved in various metabolic pathways to monitor normal function of the host cells, it is essential that Leishmania-induced dysfunction of this organelle should totally be repaired during treatment of VL. Moreover, a group of human diseases can occur when peroxisomal properties are impaired. It may lead to genetic disorder resulting in various other complications. It is thus logical to investigate whether treatment of this parasite-borne disease with the existing drugs can reverse peroxisomal defects developed due to Leishmania infection to avoid post therapeutic problems which may occur due to unavailability of specific non toxic drugs against this pathogenic disease. In the present work it has been clearly shown that resumption of normal peroxisomal function could not be attained when one of the existing drugs SAG was used for chemotherapy against VL.

2. Materials and methods

2.1 Materials

All the reagents except fetal calf serum (FCS), Medium-199 and sodium antimony gluconate (SAG) were purchased from Sigma Chemicals, USA. Fetal calf serum and Medium-199 were obtained from Gibco BRL, USA. SAG was procured from GlaxoSmithKline, UK. L. donovani strain MHOM/IN/AG/83 was obtained from Indian kala-azar patient and maintained by intracardial passage every 8 weeks in Syrian golden hamsters. Promastigotes were obtained by transforming amastigotes isolated from infected spleen and maintained in medium-199 supplemented with 10% fetal calf serum in vitro.

2.2 In vivo study with SAG in L. donovani infected hamsters

Syrian Golden hamsters (4 weeks old weighing 50-65 gm) were infected individually with freshly purified L. donovani amastigotes through cardiac route. After 30 days of infection the animals were treated with SAG (30 mg/kg body weight). Each hamster received intramuscular injections of the drugs every alternate date for 15 days. The compound was dissolved in DMSO. The final concentration of DMSO was ≤ 0.1% (v/v). Infected hamsters in the control group received 200ml of 0.1% DMSO per animal intramuscularly. Animals of all groups were sacrificed two month after administration of last treatment. The splenic parasite burden was determined from impressions smears after Giemsa staining.
parasite burden was calculated from the following formula—organ weight (mg) x number of amastigotes per cell nucleus x (2 x 10^5). All animal experiments were approved by the local body of our institute’s animal ethics committee.

2.3 Isolation of Hamster liver peroxisomes
Peroxisomes from normal and infected liver were purified according to the procedure already reported. Hamster liver was dissected out to homogenize with buffer at 4 °C containing 0.25 M sucrose, 10 mM TES of pH 7.5, 1 mM EDTA, 0.4 mM phenylmethylsulfonyl fluoride, 0.2 mM leupeptin and 0.1% ethanol. Light mitochondrial fraction from the homogenate was separated by differential centrifugation and then suspended in the same TES-EDTA buffer mentioned above at the 1:1 ratio of liver weight to buffer. 30% Nycodenz (w/v) was prepared in a buffer of pH 7.5 that contained 10 mM TES and 1 mM EDTA. 2 ml of the light mitochondrial suspension was layered over 10 ml of freshly prepared nycodenz solution and centrifuged at 105000×g for 50 min in a Sorvall-ultra 80 ultracentrifuge using T-865 fixed angle rotor. After centrifugation the interfacial materials were aspirated off to obtain pelleted peroxisome which was separated at the bottom of the centrifuge tube. Purified peroxisomes were then suspended in a minimum volume of homogenizing buffer to store at -20 °C.

2.4 Western blot analysis
Peroxisomal proteins were separated by a 10% SDS–PAGE followed by affinity transfer blotting using nitrocellulose paper.

2.5 Activity staining for SOD
Peroxisomal SOD was separated on a 10% nondenaturing polyacrylamide gel for activity staining. Gels were incubated in solution A (20 mg Nitro Blue Tetrazolium in 10 ml glass distilled water) for 20 min, then in solution B (4mg Riboflavin, 0.4 gm K_2HPO_4 and 600 µl TEMED in 50 ml glass distilled water, pH 7.8) for another 20 min and finally illuminated until white bands appeared on a blue background.

2.6 Assay of enzymes
Catalase activity was determined by monitoring the decomposition of H_2O_2 at 240 nm. SOD activity was assayed by determining ability of this enzyme to inhibit pyrogallol autoxidation rate. The assay mixture contained 0.2mMpyrogallol in air-equilibrated 50mMTris–cacodylic acid buffer, pH 8.2, and 1mM ethylenediaminetetraacetic acid. Rate of autoxidation was obtained by monitoring the increase in absorbance at 420 nm in a Hitachi spectrophotometer, No U2000. SOD has the ability to inhibit autoxidation and the extent of inhibition is taken as the measure of enzymic activity. Protein was determined using Folin and Ciocalteu’s phenol reagent.

2.7 Electron microscopy
Peroxisomes were fixed, processed to embed in Spur medium to cut thin sections, and then stained to examine under an electron microscope (Hitachi-H600) at 75KV.

2.8 Immunofluorescence
Peroxisomess were fixed with formaldehyde in PBS for 15min, permeabilized 0.3% Triton X-100 in PBS for 5min, and then blocked with 3% bovine serum albumin in PBS for 30min.
were first incubated with appropriate primary antibodies (1:100) for 30 min followed by Alexa 633 coupled secondary antibody (1:100) for another 30 min and then examined under TCS-SP Leica confocal microscope having krypton–argon mixed-laser facility.

2.9 Statistical analysis
Statistical analyses were conducted through Student’s t-test as described\textsuperscript{156}

3. Results

3.1 Effect of SAG on Leishmania infected hamsters macrophages
Following 30 days of \textit{L. donovani} infection the animals were treated SAG at a dose of 30 mg/kg body weight, dissolved in DMSO, and the splenic parasite burden was determined after two months of the last dose given as described in materials and methods. In case of the SAG treated animals, no parasite was detected (data not shown).

3.2 Electron microscopic study of Peroxisomes after SAG treatment
Peroxisomes were isolated from normal, \textit{Leishmania} infected as well as cured SAG treated hamster livers and processed for electron microscopic studies. In case of normal peroxisomes, electron dense core was observed inside the matrix and the average diameter was found to be 0.37 \(\mu\)m (Figure 1A). But peroxisomes isolated after infection were found to be swollen with severe membrane distortion, and the electron core became scattered, and a few of them had no core inside. The average diameter of the swollen organelle was determined to be 0.82 \(\mu\)m (Figure 1B). Peroxisomes obtained after SAG treatment showed interesting results. The membranes were no more distorted and the organelles were reduced almost to their normal size. The average diameter was found to be 0.4 \(\mu\)m (Figure 1C). But in most of the organelles, the dense electron core inside the matrix was not detected.

![Fig. 1. Electron micrograph of hamster liver peroxisomes (x 50,000). Peroxisomes after pre- and post-fixation with 6% gluteraldehyde and 1% osmium tetroxide were examined under Hitachi electron microscope at 75 KV. A. Normal Peroxisomes, B. Peroxisomes post 90 days infection, C. post SAG treatment](image)

3.3 Assay of peroxisomal enzymes before and after SAG treatment
Peroxisomal marker enzyme catalase was assayed in peroxisomes isolated from normal, \textit{L. donovani} infected as well as cured animal livers after SAG treatment. Specific activity of Catalase is known to be diminished during \textit{Leishmania} infection\textsuperscript{157} and the same was obtained in this case (Table 1), but catalase activity was not restored back in peroxisomes obtained from SAG treated cured hamsters (Table 1). Activity of another important peroxisomal enzyme SOD was also not detected in case of peroxisomes isolated after SAG treatment (Table 1).
Table 1. Enzyme activities in normal peroxisome, peroxisome after 90 days infection and peroxisomes after post SAG treatment. Results are in mean ± S.D. for three different experiments.

<table>
<thead>
<tr>
<th>Enzyme Specific Activity (unit/mg protein)</th>
<th>Normal</th>
<th>Infected</th>
<th>SAG Treated</th>
</tr>
</thead>
<tbody>
<tr>
<td>Catalase</td>
<td>3373</td>
<td>1884</td>
<td>1763</td>
</tr>
<tr>
<td>SOD</td>
<td>3.2</td>
<td>ND</td>
<td>ND</td>
</tr>
</tbody>
</table>

3.4 Protein profiles of peroxisomes

Figure 2 indicates that the peroxisomal protein profiles of normal, infected and SAG treated cured hamsters livers were different when subjected to SDS-PAGE. Densitometric scanning pointed to three distinct protein bands of Mr 104.5, 80.6 and 50.4, although when found to be present in peroxisomes after infection and after SAG treatment, they could not be detected in normal peroxisomes. Significant differences were also observed for the protein bands of Mr 30.2, 28.0, 20.9, 10.3 and 9.7 (Figure 2).

Fig. 2. SDS-PAGE analysis of peroxisome prepared from normal hamster liver (Lane 1), hamster liver infected with Leishmania parasite for 90 days (Lane 2), post SAG treatment (Lane 3). 40 μg protein samples were run on 10% polyacrylamide gel.

The same PAGE when subjected to Western Blotting and probed with anti Cu-Zn SOD to find out the presence of SOD in the organelles isolated at various conditions, showed that SOD was not detected in peroxisomes obtained infected as well as SAG treated cured hamsters (Figure 3).
3.5 Impact on SOD activity
Activity staining of native polyacrylamide gels clearly showed that SOD activity of infected and SAG treated peroxisomes was totally lost (Figure 4).
In a separate experiment, presence of SOD was examined using anti Cu-Zn SOD under confocal microscope. SOD was not detected in infected (Figure 5B) as well as in SAG treated (Figure 5C) hamster liver peroxisomes. Normal peroxisomes were used as positive controls (Figure 5A).

4. Discussion
Peroxisomes are abundant in host reticuloendothelial cells\textsuperscript{158}. The role and importance of peroxisomes in a wide variety of metabolic pathways, including longchain fatty acid β-oxidation, ether-linked glycerolipid biosynthesis, and \text{H}_2\text{O}_2-based respiration, have been thoroughly studied\textsuperscript{159-161}. The importance of fatty acid β-oxidation for the survival of microorganisms in other parasitic and fungal diseases has already been documented\textsuperscript{162-164}. Hamster liver peroxisomes are found to be severely damaged, both morphologically and biochemically, during \textit{Leishmania} infection\textsuperscript{157}. The structure and function of this host organelle are reported to be highly affected by the parasitic attack. The pentavalent antimonial compound SAG are widely used as first-line chemotherapeutic\textsuperscript{165, 166} agents against all forms of leishmaniasis including visceral leishmaniasis\textsuperscript{167, 168}. In this work, we have assessed the status of host liver peroxisomes after the complete treatment of the \textit{L. donovani} infected animals with standard dose of SAG.
Fig. 4. Activity staining of SOD after conducting non-denaturing PAGE. Staining was carried out according to the procedure described under Materials and methods. Normal peroxisomes (Lane 1), peroxisomes post 90 days infection (Lane 2), peroxisomes post SAG treatment (Lane 3).

Fig. 5. Confocal microscopy of Normal Peroxisomes After treatment. Anti Cu-Zn SOD peroxisomes were examined under a Leica DM IRB inverted microscope. Normal peroxisomes (Set A), peroxisomes post 90 days infection (Set B), peroxisomes post SAG treatment (Set C).
Peroxisomes were isolated by standard procedure from normal hamsters, also from animals after *Leishmania* infection and from hosts after complete cure of the parasitic disease using conventional SAG treatment. When parasite burden was measured two months after administration of the last dose, no parasite was detected in hamster spleen or liver, and that gave indication towards the complete cure of the disease.

The diameter of normal peroxisome is known to vary between 0.5 to 1.0 µm in animal tissues. However, the average diameter of hamster liver peroxisome was found to be 0.37 µm. As evidenced by electron microscopic examination the morphology of the peroxisomes after parasite infection was found to be different. Membrane structure was damaged and the dense electron matrix was unevenly distributed, with the appearance of dark patches having an average diameter of 0.82 µm. The presence of large peroxisomes suggests the swelling of these organelles during infection. But the average diameter of peroxisomes isolated from hamsters after SAG treatment was determined to be 0.4 µm, which is quite similar to that of the normal organelles, indication that damaged peroxisomes are morphologically repaired after SAG treatment.

On the other hand, the key peroxisomal enzyme catalase was found to be less active after host-parasite interaction. But when catalase activity was examined in SAG treated hamster liver peroxisomes, it was found that the enzyme activity was not restored back, even after complete cure of the parasitic disease, indicating that the peroxisomes are biochemically not repaired after SAG treatment.

SOD is one of the key enzymes of oxygen defense system. It is known to be an essential factor in mediating normal cellular functions. It is already reported that peroxisomal SOD activity is not detected after *Leishmania* infection. In this study, the results clearly revealed that this essential enzyme is absent even in the peroxisomes isolated from hamster livers, after SAG treatment. In a separate experiment, Western Blot analysis showed no band for SOD in case SAG treated host liver peroxisomes. Activity staining of native polyacrylamide gels also showed that activity of peroxisomal SOD, was lost completely after parasitic disease, and could not be detected after SAG treatment. Confocal microscopic studies also revealed the same results, indicating that peroxisomal SOD activity was not restored back post SAG therapy.

Difference in protein profile in peroxisomes isolated from normal, infected as well as SAG treated hamster livers also indicate towards the status of peroxisomes. Some distinct bands were found in the infected and SAG treated peroxisomes, but were absent in normal peroxisomes. During *Leishmania* infection, some proteins get damaged and also proteolytic cleavage takes place, as result, more bands were obtained in infected peroxisomes. The similarity in protein profiles between infected and SAG treated peroxisomes revealed that the biochemical properties of peroxisomes are not restored properly after SAG treatment.

Considering the knowledge we have gathered from the present work, it is clearly established that the peroxisomes, that get damaged during *Leishmania* infection are morphologically repaired after treatment with a common antileishmanial agent SAG, but the biochemical properties are not restored back.

PKDL is a common problem in kala azar patients but the actual reason behind the occurrence of this disease is not yet known. Several peroxosomal disorders like ZS, IRD etc are also reported. Treatment with a common drug SAG, could repair the morphology of the distorted peroxisomes but the biochemical properties are not restored.

We propose that the peroxisomal defects caused by infection of the parasite *Leishmania donovani* lead to induce symptoms similar to genetic defects in the biogenesis of
peroxisomes. Although there is no authentic evidence available at this time, defective peroxisomes may also be the root to crop up PKDL in post kala-azar patients. Extensive investigations are still needed to sort out various unresolved problems in these areas to provide clues for new drug development against leishmaniasis. Present research in our laboratory is oriented towards this direction.

5. Conclusion

VL is a deadly disease of viscera caused by the parasite *Leishmania donovani* and also known as kaka-azar. The internal organs particularly liver, spleen, bone marrow and lymph nodes are attacked when the parasitic protozoa is transmitted to humans by infected female sand fly bites. People of Mediterranean and adjacent countries are attacked with this pathogen in the form of flagellated promastigotes. However, aflagellated amastigotes are found inside and outside of the affected reticuloendothelial cells.

At this time there is no definite antileishmania agent which may be used confidently to treat VL for a permanent cure. Drugs of choice available in the market, are being taken up selectively by trial and error methods. Most of them are too toxic to cause severe secondary infections and at the same time very costly to afford. Moreover, drug resistance is one of the acute problems to provide guarantee for a complete cure of VL. As a result there is an urgent need to explore for new drug development against this parasitic disease which is lethal, if untreated.

While exploring our attempts to identify an authentic chemotherapeutic agent against VL we also took into account the occurrence of PKDL. There are reports to believe that cured kala-azar patients as certified by the physicians, are susceptible to return for the treatment of PKDL.

Cause of PKDL is still not known. It is possible that the said pathogenic disease is occurred due to continued impaired function of the internal host organs even after treatment with commonly found drugs. We have shown that host liver peroxisomal properties are impaired due to the attack by *Leishmania* pathogen. We have also reported that the affected peroxisomes are not repaired after adequate treatment with the existing drugs. Treated animals with deficiency in peroxisomal properties also do not survive for a long time.

It is known that that peroxisomal defects lead to various genetic disorders. It is still not known whether in treated patients of VL who are likely to carry peroxisomal deficiencies even after traditional treatments, symptoms similar to genetic defects in the biogenesis of peroxisomes are developed.

A novel chemotherapeutic agent has been discovered in our laboratory to prove that this antileishmania agent has the ability to repair peroxisomal damage which takes place due to attack by VL. This finding may lead to provide effective clues to develop new and more potent drugs to stop recurrence of this pathogenic disease in other forms including chance to trigger symptoms develop due to peroxisomal disorder.

6. References


Repair of Impaired Host Peroxisomal Properties Cropped Up Due to Visceral Leishmaniasis May Lead to Overcome Peroxisome Related Genetic Disorder Which May Develop Later...


Repair of Impaired Host Peroxisomal Properties Cropped Up Due to Visceral Leishmaniasis May Lead to Overcome Peroxisome Related Genetic Disorder Which May Develop Later...


[107] de Duve, Ch. The separation and characterization of subcellular particles. Harvey Lect. 1965 59: 49-87


Repair of Impaired Host Peroxisomal Properties Cropped Up Due to Visceral Leishmaniasis May Lead to Overcome Peroxisome Related Genetic Disorder Which May Develop Later...


[117] Lazarow, P.B. and de Duve, C. A fatty acyl-CoA oxidizing system in rat liver peroxisomes; enhancement by clofibrate, a hypolipidemic drug. Proc Natl Acad Sci USA.1976 73: 2043-60

[118] Demoz, A., Daniel, K., Lie, A.O. and Berge, R. K. Modulation of plasma and hepatic oxidative status and changes in plasma lipid profile by n-3 (EPA and DHA), n-6 (corn oil) and a 3-thia fatty acids in rats. Biochimica et Biophysica Acta. 1994 1199(3): 238-244


[125] Raymon, G.V. Peroxisomal Disorders, Current Opinions in pediatrics, 1999 11: 572-76


Repair of Impaired Host Peroxisomal Properties Cropped Up Due to Visceral Leishmaniasis May Lead to Overcome Peroxisome Related Genetic Disorder Which May Develop Later...


The studies on genetic disorders have been rapidly advancing in recent years as to be able to understand the reasons why genetic disorders are caused. The first Section of this volume provides readers with background and several methodologies for understanding genetic disorders. Genetic defects, diagnoses and treatments of the respective unifactorial and multifactorial genetic disorders are reviewed in the second and third Sections. Certainly, it is quite difficult or almost impossible to cure a genetic disorder fundamentally at the present time. However, our knowledge of genetic functions has rapidly accumulated since the double-stranded structure of DNA was discovered by Watson and Crick in 1956. Therefore, nowadays it is possible to understand the reasons why genetic disorders are caused. It is probable that the knowledge of genetic disorders described in this book will lead to the discovery of an epoch of new medical treatment and relieve human beings from the genetic disorders of the future.

How to reference
In order to correctly reference this scholarly work, feel free to copy and paste the following: