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Retrospective Analysis of Leishmaniasis in Central Tunisia: An Update on Emerging Epidemiological Trends

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1. Introduction

This study aimed at describing the spatio-temporal distribution of leishmaniasis in patients who have sought diagnosis in the laboratory of Parasitology of Farhat Hached hospital, Sousse, Tunisia, across the 1986-2010 period in order to: i) highlight important features and trends of leishmaniasis and its epidemiology; ii) and to assess whether the activity of the unit reflects the situation of the disease at the national level and whether it could constitute an indicator of public health relevance.

2. Current situation of leishmaniasis in Tunisia

Tunisia is located in Northern Africa, bordering the Mediterranean sea between Algeria and Libya. Its climate is of the hot temperate Mediterranean type. Bioclimatic zones range from humid in the north to saharian in the extreme south (Figure 1). The country is divided into 24 governorates, each composed of a variable number of “delegations”; and each delegation is subdivided into localities also named “imadas” (Figure 2). Four forms of leishmaniasis are known to occur in Tunisia: i) the sporadic cutaneous leishmaniasis (SCL); ii) the chronic cutaneous leishmaniasis (CCL); iii) the zoonotic cutaneous leishmaniasis (ZCL); iv) the visceral leishmaniasis (VL).

2.1 The sporadic cutaneous leishmaniasis (SCL)

This form is caused by dermatropic zymodemes of *Leishmania infantum*, mainly the zymodeme MON 24 and to a lesser extent MON 1 (Aoun et al., 2000, 2008; Ben Ismail et al., 1986, 1992; Gramiccia et al., 1991; Haouas et al., 2007; Kallel et al., 2005, 2008b). The dog is supposed to be the reservoir as it is the case for the visceralotropic zymodemes and *Phlebotomus perfiliewi* as the phlebotomine vector. But other vectors can not be excluded; following an outbreak of SCL in the locality of Oued Souani, le Kef governorate, *P. langeroni* was found infected by *L. infantum* (Guerbouj et al., 2007).
SCL was first described in 1917 in a patient from Sakiet Sidi Youssef (Le Kef governorate) located next to the Algerian frontier (Nicolle & Blanc, 1917), and again in 1945 (Chadli et al., 1968; Vermeil, 1956). Since 1945, the disease has sporadically been reported with, however, an incidence that gradually increased from a median of one case/year between 1945 and 1955 to 22 cases/year in the 1990s and 59 cases/year in the 2000s (Table 1; Anonymous; Ben Abda et al., 2009; Ben Said et al., 2006; Ben Ismail & Ben Rachid, 1989; Ben Rachid et al., 1983, 1992; Chadli et al., 1968; Chaffai et al., 1988; Ladjimi & Lakhoua, 1955; Vermeil, 1956). SCL has long been supposed to be confined to the humid and subhumid bioclimatic areas north to the “dorsale” or “tunisian Ridge” (the eastern extension of the Atlas mountains), in rural areas where its distribution overlaps with that of VL (Ben Ismail et al., 1989; Ben Rachid et al., 1983, 1986; Chadli et al., 1986; Vermeil, 1956). However, over the last three decades, many cases originating from central Tunisia governorates (Monastir, Sousse, Mahdia and Kairouan) and referred to our laboratory were found to be very suggestive of SCL on the basis of epidemiological, clinical and parasitological criteria. Typed strains obtained from some of these patients proved indeed to be L. infantum (Ben Said et al., 2006). These data were very indicative of SCL spread towards the south of the country. These findings were further confirmed by additional reports (Aoun et al., 2000, 2008; Ben Abda et al., 2009; Kallel et al., 2008a, 2008b). The revised SCL distribution is shown in figure 3. SCL usually occurs sporadically within no particular season. However, over the last few years, local outbreaks were reported like that of Sidi Bourouis in Siliana governorate, which involved more than 30 patients over a short period (Bel Hadj et al., 2003).
Fig. 2. Map of Tunisia showing its subdivision into 24 governorates (Grand Tunis is composed of 4 governorates).

<table>
<thead>
<tr>
<th>Years</th>
<th>Nb of cases</th>
<th>Median</th>
</tr>
</thead>
<tbody>
<tr>
<td>1945-55</td>
<td>10</td>
<td>1.0</td>
</tr>
<tr>
<td>1956-67</td>
<td>17</td>
<td>1.4</td>
</tr>
<tr>
<td>1968-78</td>
<td>84</td>
<td>8.4</td>
</tr>
<tr>
<td>1979-89</td>
<td>151</td>
<td>15.1</td>
</tr>
<tr>
<td>1990-99</td>
<td>217</td>
<td>21.7</td>
</tr>
<tr>
<td>2000-10</td>
<td>652</td>
<td>59.3</td>
</tr>
<tr>
<td>Total</td>
<td>1131</td>
<td>17.1</td>
</tr>
</tbody>
</table>

Table 1. Number of recorded sporadic cutaneous leishmaniasis cases since 1945.

Clinically, SCL is characterized in more than 90% of patients by a small single ulcerated or lupoid lesion of the face that often lasts longer than one year and up to three years (Bel Hadj et al., 1996; Ben Ismail & Ben Rachid, 1989; Chaffai et al., 1988; Masmoudi et al., 2007). On a parasitological level, amastigotes of *L. infantum* are smaller than those of *L. major*, usually...
less than 3μ, and promastigotes are difficult to maintain on NNN medium (Aoun et al., 2000, 2003).

Fig. 3. Updated distribution of sporadic cutaneous leishmaniasis in Tunisia.

2.2 The chronic cutaneous leishmaniasis (CCL)
Formerly known as anthroponotic CL, CCL sporadically occurs in micro-foci located in the south-eastern presaharian and saharian areas of Tunisia. Its distribution is rural, sub-urban and urban; and cases are reported from houses, farms, and even from troglodytes of Tataouine (Tataouine governorate) and Matmata (Gabès governorate), with a median
annual incidence of 10 cases/year (Figure 4; Ben Ismail & Ben Rachid, 1989; Ben Rachid et al., 1983, 1992; Chadli et al., 1968). CCL was first reported in 1957 and nearly nothing was known about the disease before this date. In 1979, an outbreak involving 47 individuals arose in Tataouine; and the epidemiological investigation, which included the isoenzymatic typing of strains isolated from the patients, led to the identification of the parasite as the MON 8 zymodeme of *L. tropica*, named *L. killicki*, a species previously described in Kenya, Namibia, Yemen and, more recently in Algeria and Libya (Harrat et al., 2009; Rioux et al., 1986; Sang et al., 1994).

Surprisingly, over the last decade, cases of CCL were reported in patients originating from areas where this form has never been previously reported, the first case being in a child from Meknassy in Sidi Bouzid governorate (Haouas et al., 2005). Later, additional cases were reported from Kairouan, Gafsa and Siliana governorates, very far from the classical foci of Tataouine (Aoun et al., 2008; Ben Abda et al., 2009; Bouratbine et al., 2005).
Hence the actual distribution and incidence of CCL need further investigations and obviously should be revised.

*L. killicki* is transmitted by *Phlebotomus sergenti*, but the reservoir is still debated. Median age of patients suffering of CCL is 21 years (Ben Abda et al., 2009; Ben Ismail & Ben Rachid, 1989). Clinically, CCL most often presents as single or very few lesions on the face or limbs, that are dry, extensive, and chronic, lasting for up to six years (Ben Abda et al., 2009; Ben Ismail & Ben Rachid, 1989; Chaffai et al., 1988; Masmoudi et al., 2007).

### 2.3 The zoonotic cutaneous leishmaniasis (ZCL)

ZCL is by far the most frequent and the most widely distributed form of CL in Tunisia where it constitutes a major public health problem. It is endemo-epidemic in endemic areas of central and southern Tunisia (Figure 5). It is caused by *L. major* and transmitted by the zoo-anthropophilic *Phlebotomus papatasi* sandfly which is mainly encountered and caught in and around the rodent burrows, and less in and around human habitations (Ben Ismail et al., 1987b, 1987c; Ben Rachid et al., 1992; Ghrab et al., 2006; Helal et al., 1987). The reservoirs are rodents of the *Psammomys* and *Meriones* genera. The main one is *P. obesus*, a prolific diurnal rodent that is very abundant in arid and subsaharian areas. Its feeding requirements consist exclusively of chenopodiaceae (*Salicornia*, *Salsola*, *Atriplex*) that mainly grow in sandy, humid and salty soils unsuitable for agricultural purposes (Ben Ismail et al., 1987a; Ben Rachid et al., 1992; Fichet-Calvet et al., 2003). *Psammomys* infection rate may reach 100%. The hygrophilic nocturnal rodents, *Meriones shawi* and *Meriones libycus* act as secondary reservoirs and are responsible of the spread of the disease because of their migratory habits. *Meriones* are granivorous and build their burrows in jujube trees (*Zizyphus*) surrounding cereal fields and often cause important agricultural damage (Ben Ismail & Ben Rachid, 1989; Ben Rachid et al., 1992). All *Leishmania* strains isolated so far from humans, rodents and phlebotomine vectors are of the zymodeme MON 25 (Aoun et al., 2008; Ben Abda et al., 2009; Ben Ismail et al., 1986; Haouas et al., 2007).

ZCL was first described in 1882 in and around Gafsa oases, and termed “clou de Gafsa” (Deperet & Bobinet, 1884). From this date and up to the beginning of the 20th century, many additional outbreaks were reported in the same area. Then, the disease continued to occur on a very sporadic mode, and nearly disappeared (Chadli et al., 1968; Vermeil, 1956).

In 1982, a large outbreak arose in Nasrallah delegation (Kairouan governorate), near to the recently completed Sidi-Saad dam (Ben Ammar et al., 1984). From there, the disease rapidly spread to cover large rural and sub-urban parts of the central and south-western neighbouring governorates, so that, by 1986, 10 governorates were involved (Figure 6). In season 1991-1992, ZCL extended further south-east to Medenine and Tataouine governorates. All along the outbreak, Sidi Bouzid, Gafsa and Kairouan governorates have remained the leading areas in terms of incidence (Anonymous). However, from the early 1990s and up to date, Tataouine, Tozeur, Médénine, Kébili, Gabès, Sfax and Kasserine have emerged as active and stable foci (Figure 5). In the last few years, some level of ZCL spread towards the north (Siliana, Béja, Le Kef, Tunis and Zaghouan governorates) was registered, which is somewhat surprising (Ben Abda et al., 2009).

The number of annual recorded cases rapidly grew from 182 cases in 1982 to > 18000 cases by 1987, > 65000 cases by 1999. Up to date, > 120000 cases were reported and the epidemic is still going on. It should be mentioned however, that the actual number of cases is supposed to be underestimated and would exceed 150000 cases (Chahed et al., 2002). The number of annual recorded cases greatly varied and ranged from 1129 cases in 1995 to > 15000 in 2004,
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Fig. 5. Distribution of zoonotic cutaneous leishmaniasis according to incidence (expressed as cases/100,000).

with an average of 4700 cases/year (Figure 7). The annual incidence showed the same fluctuations. It ranged between 30/100,000 in the less affected areas and > 500 in Kairouan, and > 1000 in Gafsa and Sidi Bouzid in 2004 (Anonymous). These fluctuations are at least in part related to the dynamics of rodents’ populations. It was demonstrated that the distribution of *P. obesus* is of paramount importance in the epidemiology of the human disease (Ben Ismail et al., 1987a; Ben Ismail & Ben Rachid, 1989; Ben Salah et al., 2007; Fichet-Calvet et al., 2003). On the other hand, control programmes are undoubtedly very effective and beneficial in terms of incidence. Indeed, a rodent control project, consisting of ploughing lands with growing chenopods and reafforestation with acacia and other plants was launched in 1992 in Sidi Bouzid city and suburbs that resulted in a dramatic decrease in the incidence of the disease in the area; but the intensity of the epidemic grew again as soon as the control programme stopped (Figure 7). From the mid 2000s, similar control measures have been carried out again in Sidi Bouzid and then in Sidi El Heni...
delegation which is the main transmission focus in Sousse governorate, and led to an obvious decrease of incidence in both foci. The ongoing outbreak that started in 1982 near the Sidi-Saad dam may be explained by the interruption, as a result of the construction of the dam, in the flooding that frequently occurred in the area and used to decimate a high proportion of rodents every year. In addition, the enrichment of the area’s ground water helped the chenopodiaceae to grow abundantly, thereby increasing the food source of *Psammomys*. On the other hand, *Atriplex*, a plant grown in large quantities as a sheep fodder is much appreciated by *Psammomys*. Furthermore, humidity created by the dam is highly suitable for the sandflies.

Fig. 6. Spread of zoonotic cutaneous leishmaniasis during the period 1982-1986 (Ben Ismail R., unpublished).

Fig. 7. Annual number of reported zoonotic cutaneous leishmaniasis cases at the national level from 1982 up to 2010.
One of the characteristics of ZCL is its marked seasonal occurrence as most cases are observed between October and January (Ben Ismail & Ben Rachid, 1989; Ben Rachid et al., 1992). Clinically, ZCL most often presents as multiple, inflammatory ulcers on the face and the limbs that usually scar in less than 6-8 months, and affects all ages with a median of 24 years (Ben Abda et al., 2009; Ben Ismail & Ben Rachid, 1989; Chaffai et al., 1988; Masmoudi et al., 2007; Zakraoui et al., 1995).

2.4 Visceral leishmaniasis (VL)

VL has been known to sporadically occur in Tunisia, since 1903 where the first case of Mediterranean VL was described in a child living in the suburbs of Tunis. The disease is caused by *Leishmania infantum*, mostly zymodeme MON 1 and to a lesser extent zymodemes MON 24 and MON 80 (Aoun et al., 2001, 2008; Bel Hadj et al., 1996, 2000, 2002; Ben Ismail et al., 1986; Haouas et al., 2005; Kallel et al., 2008b). It is transmitted by *Phlebotomus perniciosus* sandfly, and the dog is the exclusive reservoir, with an infection prevalence rate ranging from 5% to 26% (Ben Ismail et al., 1986; Ben Rachid et al., 1992; Bouratbine et al., 1998).

Since 1903, 2449 VL cases were reported in Tunisia (Table 2).

<table>
<thead>
<tr>
<th>Period</th>
<th>Nb cases</th>
<th>Median/year</th>
</tr>
</thead>
<tbody>
<tr>
<td>1903-1956</td>
<td>151</td>
<td>2.8</td>
</tr>
<tr>
<td>1957-1967</td>
<td>142</td>
<td>12.9</td>
</tr>
<tr>
<td>1968-1981</td>
<td>178</td>
<td>12.7</td>
</tr>
<tr>
<td>1982-1989</td>
<td>578</td>
<td>72.3</td>
</tr>
<tr>
<td>1990-2000</td>
<td>720</td>
<td>65.5</td>
</tr>
<tr>
<td>2001-2010</td>
<td>680</td>
<td>68.0</td>
</tr>
<tr>
<td><strong>Total</strong></td>
<td><strong>2449</strong></td>
<td><strong>22.7</strong></td>
</tr>
</tbody>
</table>

Table 2. Number of visceral leishmaniasis cases reported in Tunisia (1903-2010)

Up to 1981, incidence was low to moderate and nearly all cases were reported from Zaghrouan, North-West (Le Kef, Béja, Jendouba, Siliana), Tunis and Sousse governorates, located in the humid, sub-humid and semi-arid zones (Anderson et al., 1934, 1938; Ben Ismail et al., 1986; Ben Rachid et al., 1983; Chadli et al., 1968; Khalidi et al., 1991; Nicolle, 1912; Vermell, 1956). From the early 1980s, the incidence markedly increased and the disease progressed towards the south, mainly to Kairouan governorate and to a lesser extent to Sfax, Sidi Bouzid, Kasserine and Tozeur governorates, together with an increase in canine leishmaniasis (Ayadi et al., 1991; Bel Hadj et al., 1996; Ben Salah et al., 2000; Besbès et al., 1994; Bouratbine et al., 1998; Chargui et al., 2007; Pousse et al., 1995). Indeed, from the 1980s, the region of Kairouan emerged as a highly active VL focus, and by 1991-1992 it was recognized as the most active one, with 30 to 55% of reported cases (Anonymous).

The distribution of VL is given in figure 8. Highest number of cases was reported in 1922 (n = 130), 1993 and 2006 (n = 122), and 2005 (n = 120). It is worth mentioning that VL has nearly disappeared between 1974 and 1980, as a result of the anti-malaria campaign which included extensive insecticide spraying (Ben Rachid et al., 1983). The disease sporadically occurs in rural and to a lesser extent in suburban areas and mainly affects children under five years, cases in immunocompetent adults being less than 5% (Ben Ismail & Ben Rachid, 1989; Ben Rachid et al., 1983; Besbès et al., 1994; Bouratbine et al., 1998; Hammoud et al.,...
In HIV Tunisian patients, VL is infrequent; it is best diagnosed by PCR (Kallel et al., 2007).

3. Materials and methods

The laboratory of parasitology of Sousse, the third major Tunisian city which is located in the central, coastal part of Tunisia, was first created in 1986. It can be considered as a central laboratory ensuring the diagnosis of parasitic diseases’ cases including leishmaniasis, sent by generalists, dermatologists, pediatricians or hematologists either from the same hospital or other hospitals mainly from governorates of central Tunisia like Kairouan, Sidi Bouzid and Mahdia. Results are usually sent back to the referring doctors so the patients can be managed as appropriate. Since its creation, the laboratory has been involved in the diagnosis of all forms of leishmaniasis.

For purposes of this retrospective study, VL and CL patients’ data, including age, sex, geographical origin and the likely place of contamination, clinical presentation and evolution without or after treatment were collected.

3.1 Diagnosis of cutaneous leishmaniasis

The diagnosis of CL is usually achieved by the demonstration of amastigotes in Giemsa-stained smears from the fluid obtained by scraping the edges of the cutaneous lesion with a
sterile lancet. In addition, the size of the identified *Leishmania* is carefully evaluated, because small amastigotes (< 3 μ) are evocative of *L. infantum* and *L. killicki* rather than *L. major* whose amastigotes are larger. Over the last decade, lesions evocative of CL but found negative in direct examination were further submitted to PCR technique, known to be more sensitive.

### 3.2 Diagnosis of visceral leishmaniasis

#### 3.2.1 Parasitological diagnosis

Diagnosis of VL is usually based on the demonstration of amastigotes in Giemsa stained bone-marrow aspirates. However, in our laboratory, bone-marrow smears are mainly carried out and read in doubtful cases, because they usually are performed and interpreted by the haematologists and sometimes the pediatricians themselves and the results of the examination are later sent to us together with sera specimens.

For the purpose of parasitological diagnosis of VL, since 1996, we have been using the cytoconcentration of peripheral blood according to the technique of Petithory et al, 1997. Cytoconcentration is usually practiced first, and when positive the painful bone marrow aspiration is no more needed and the diagnosis of VL is established.

#### 3.2.2 Serodiagnosis of leishmaniasis

All sera from suspected or confirmed VL cases are sent to our laboratory for serodiagnosis. Our reference technique is the fluorescent antibody test (FAT). It is carried out by using *Leishmania*-spot IF slides (bioMerieux, France) on sera serially diluted starting from 1/100. Sera showing fluorescence at ≥ 1/200 dilution are considered positive.

In addition to FAT, we have been using, since 2003, the rK39 dipstick test (DST). The first one we used was from Inbios International, USA (kalazar detect®). Then, since 2007 and up to date, we have been using the Dia.Med IT Leish® (Dia Med, Switzerland), according to Saghrouni et al., 2009.

### 3.3 Culture of *Leishmania*

Our laboratory has mainly been involved in the culture of dermotropic *Leishmania*, even though isolates from VL bone-marrow aspirates were occasionally cultured. Culture was mainly carried out for the purpose of typing strains by isoenzyme electrophoresis (IEE) technique. Media we have been using are the NNN (Nicolle Novy Mc Neal), the BHI (brain heart infusion agar) and the CRS (coagulated rabbit serum), the last being best for cutaneotropic *L. infantum* strains. *Leishmania* isolates were typed either in the IEE unit of the LEEP or in the laboratory of parasitology of the Faculty of Pharmacy, Monastir, where an IEE unit was set up in 2001.

### 3.4 Molecular biology

Up to 2008, our laboratory was not equipped for molecular biology techniques. In 2007, as part of a collaborative project on “molecular tools for accurate diagnosis and better assessment of leishmaniases”, codirected by the LEEP and our own laboratory and financed by the IAEA, a molecular biology unit was set up in our department and has just started working. In the meanwhile we have been collaborating with the molecular biology unit of the LEEP, mainly for identifying *Leishmania* strains. The first study, conducted in 2005, aimed at identifying strains isolated from patients suspected for having SCL, by using K-DNA-PCR according to Smyth et al., 1992 and PCR-RFLP according to Guerbouj et al., 2001 (Ben Said et al., 2006).
Later, additional strains isolated from CL patients were typed by a novel Multiplex PCR (Saadi et al., in preparation) as part of the IAEA project that is still ongoing.

In addition, specimens from patients with lesions highly evocative of CL but found negative in parasitological examination were addressed for PCR to the laboratory of parasitology of the Faculty of Pharmacy, Monastir, where a molecular biology unit has just been set up. PCR was performed according to Chargui et al., 2005.

4. Results

4.1 Cutaneous leishmaniasis

Over the 25 year period study, 4329 patients were investigated for CL. Most of them were referred to our laboratory by the service of Dermatology of Farhat Hached hospital, Sousse. Leishmania parasites were demonstrated in 2087 cases (48.2%). In addition, out of 86 PCR performed on samples found negative in direct examination of dermal Giemsa stained smears, 17 were positive. So, the total number of CL cases diagnosed during the study period was 2104. Most of them were diagnosed during the last decade.

Out of the 2104 diagnosed cases 50 were confirmed or very likely SCL form. Fourteen came from areas known to be endemic for SCL (Le Kef, Jendouba, Siliana, Zaghouan and Bizerte governorates). The remaining 36 cases were from areas located in central Tunisian governorates where SCL has never been described: 13 were from Monastir, 12 from Sousse, 6 from Mahdia and 5 from Kairouan. Out of the 36 cases, 13 originated from areas known to be endemic for ZCL, in Kairouan, Mahdia and Sousse governorates. The age of patients ranged from 5.5 to 63 years (median = 28.5 years). Twenty six were males and 24 were females (sex ratio M/F = 1.08).

In three patients, the isolate proved to be L. killicki. The first patient was a 5 year old child from Meknassi in Sidi Bouzid governorate where CCL was unknown. The second was a 30 year old woman from Ghomrassen, known to be endemic for CCL. The third patient was a 21 year old woman who came from Nasrallah, one of the most active ZCL foci in Kairouan governorate and from where no CCL cases were reported before.

All the remaining 2051 patients were suffering from ZCL. 1182 were females and 869 were males (sex ratio F/M = 1.36). Their age ranged from 1 month to 90 years (median = 28 years). The annual distribution of ZCL cases diagnosed over the 25 year period is shown in figure 9. The number of cases ranged from 5 in 1997 to 443 in 2004.

The place of contamination could be ascertained in 1873 out of the 2051 patients. In the 178 remaining cases, the geographical origin of contamination could not be determined with certainty because of multiple displacements of patients across two, three or more endemic areas. In addition, some patients were originating from Libya and Algeria and were not included in the analysis of the spatio-temporal distribution of ZCL cases.

The distribution of the 1873 ZCL cases according to the area where the contamination took place is given in table 3. Most of the patients came from Sidi Bouzid (610 = 32.6%), Mahdia (494 = 26.4%), Kairouan (369 = 19.7%), and Sousse (306 = 16.3%) governorates.

The distribution of cases according to delegations inside the four governorates mentioned above is shown in figure 10, and the annual distribution of diagnosed cases in figure 11. According to seasonal distribution 1745 (85.1 %) cases were diagnosed between October and February. All confirmed CL patients were treated with local or parenteral N-methylglucamine antimoniate (glucantime®) together with cryotherapy in those with few lesions. Most of treated patients responded well to antimonial treatment, and scarring of
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Fig. 9. Annual distribution of 2051 zoonotic cutaneous leishmaniasis cases (1986-2010).

Table 3. Distribution of 1873 zoonotic cutaneous leishmaniasis cases according to governorate and delegation.
Fig. 10. Distribution of 1779 zoonotic cutaneous leishmaniasis cases diagnosed over the 25 year period originating from Sidi Bouzid, Mahdia, Kairouan and Sousse. The dots represent the number of cases. The limits of the governorates are shown in bold lines. The administrative district subdivisions are illustrated by different colours in each governorate.

Lesions was obtained in a few weeks (one to three) after the treatment was initiated. However, in some patients, the outcome was unexpectedly atypical in that the lesions took much more delay to heal as demonstrated by the persistence of *Leishmania* in direct examination. In some adequately treated patients, the lesions persisted longer than one year and up to 4.5 years in one of them. In another patient, nearly 100 glucantime® injections were needed before the lesions resolved. In some additional cases, new lesions appeared while the patient was under specific treatment for previous ulcers. On the other hand, in many patients treated with *in situ* antimonial infiltrations, sporotrichoid nodules developed a few days or weeks later, next to the treated lesion.
4.2 Visceral leishmaniasis

Up to 2010, 2784 serum samples were addressed to our laboratory for suspected VL cases. Anti-Leishmania antibodies were demonstrated in 944 sera. FAT and rK39 DST were positive in 944 (99.2%) and 822 (87.1%) cases respectively. In all positive patients, VL was confirmed by the demonstration of amastigotes in bone marrow and/or cytocentrifuged blood and/or favourable outcome under antimonial treatment. In 8 confirmed VL cases, both serological tests were negative. Thus, the total number of VL cases diagnosed over the 25 year period is 952. Three hundred and eight blood samples were investigated by the cytoconcentration technique. Among them, 201 were from confirmed VL patients, and amastigotes were demonstrated in 88 (43.7%) of them. In six cases, bone marrow examination was also negative. Apart from the confirmed VL FAT negative cases, all the remaining negative sera were from patients suffering from diseases other than VL, mainly leukaemia and various haematological disorders. Nine hundred and thirty four of the VL patients were children, aged 2 months to 14 years (median: 26.8 months). Eighteen were adults aged 19 to 39 years (median: 25.9 years); one of them was a HIV+ woman. The number of annually diagnosed cases ranged from 14 cases in 2004 to 83 cases in 2005. The highest proportion of VL cases (60%) was from Kairouan governorate, known to include the most active foci of VL in Tunisia, followed by Sousse governorate and, to a lesser extent from other areas in central Tunisia (Mahdia, Monastir, Kasserine).

All confirmed VL patients were treated with glucantime®, usually a single 21 to 28 day course. The outcome was favourable in the majority of them. Death occurred in 18 (1.9%) patients. All were children. In 4 of them, no response to antimonials was observed despite an adequate treatment regimen. They were assumed to be resistant to antimonials even
though the resistance could not be assessed by *in vitro* testing of the strain. In an additional child, initially treated by three successive courses of glucantime®, healing was ultimately obtained with ambisome® (liposomal amphotericin B).

### 4.3 Typing of *Leishmania* strains

Over the 25 year study period 103 *Leishmania* strains were typed by either IEE or PCR or both. More than 150 isolates were *in vitro* cultured. Fifteen of them were from bone-marrow aspirates in VL cases. All were typed by IEE and found to be *L. infantum*. Most cultured isolates were from CL patients. Eighty one of them could be maintained in culture media, cryopreserved and typed by IEE. The remaining isolates were either lost after repeated subculture or contaminated during sampling or subculturing. It should be mentioned that strains addressed to the laboratory of Monastir, were typed and identified at the zymodeme level, whereas those typed in the LEEP were only identified at the species level.

Sixty four isolates were typed by molecular techniques: 4 were typed by K-DNA-PCR and PCR-RFLP, and 60 by multiplex PCR. Overall, 88 CL strains were typed by either IEE or PCR technique: 7 were *L. infantum*, 3 were *L. killicki* and 78 were *L. major*. All *L. major* strains that were typed at the zymodeme level were found to be *L. major* MON 25.

### 5. Discussion

One of the main characteristics of the four leishmaniasis forms occurring in Tunisia is their changing epidemiological pattern, which has continuously been described over the last 30 years. These changes mainly consisted of the spread of all forms of the disease from their initial endemic foci towards neighbouring or farther areas.

The VL which has long been confined to the northern parts of Tunisia showed a noticeable spread towards the central and southern parts of the country, where the disease has never been previously reported. The Kairouan governorate where VL was unknown till 1980, has progressively emerged as a highly endemic area and by the early 1990s, it became the most active focus in terms of incidence (Anonymous; Bel Hadj et al., 1996; Besbès et al., 1994; Bourattine et al., 1998). These findings contrast with earlier periods where VL was mainly reported from Zaghouan, North-western and Tunis governorates and to a lesser extent from the central coastal region (Sousse) (Anderson et al., 1938; Ben Ismail et al., 1986; Ben Rachid et al., 1983; Chadli et al., 1968). In the same time, VL progressively spread further south to southern and central western governorates (Sfax, Sidi Bouzid, Kasserine, Tozeur) (Ayadi et al., 1991; Chargui et al., 2007). While spreading, the VL incidence markedly increased from a median of 12 to 68 annual recorded cases, 50 % of them being from Kairouan foci.

Similar trends were described in SCL which, up to 1980, was very rarely reported south to the Tunisian ridge (“dorsale”). From the early 1980s, and up to date, SCL has continuously been reported from central and southern parts of the country (Aoun et al., 2000, 2008; Ben Said et al., 2006; Kallel et al., 2005, 2008a).

ZCL was first reported on the end of the 19th century causing large outbreaks in Gafsa and neighbouring areas. Later, it has nearly disappeared for half a century; but reappeared again in 1982 causing a widespread epidemic that is still going on and constitutes a major public health problem in extended areas of the country, with up to date more than 150000 cases (Anonymous; Ben Abda et al., 2009; Ben Ismail & Ben Rachid, 1989).
CCL, known to be confined to limited areas in the presaharian region of Tataouine, has itself shown spread towards areas far from its original endemic foci, up to Kairouan governorate (Aoun et al., 2008; Haouas et al., 2005). The described changes are supposed to mainly originate in the agricultural development and the ecological transformations that occurred in the concerned areas and led to a marked increase in irrigated surfaces that in turn helped the reservoirs (mainly rodents) and vectors to abundantly proliferate, and created biotopes very suitable for the Leishmania cycles to establish and amplify (Aoun et al., 2008; Ben Abda et al., 2009; Ben Salah et al., 2007). In addition, urbanization of previously rural areas made the populations’ access to medical facilities easier, and consequently more and more leishmaniasis cases could be detected and diagnosed. The impact of climate change and variability on leishmaniasis (due to el Niño) was shown in Colombia and is associated to shifts in insect and animal distributions (Cardenas et al., 2008). This phenomenon was also shown to impact on the incidence of VL in Brazil and models were established to predict high risk years for VL (Franke et al., 2002). In Tunisia, the relationship between climate variability and leishmaniasis is not well studied. However, incidence rate of VL was shown to be positively correlated with mean yearly rainfall and continentality index; a rainy year is followed 2 years later by an increase in VL cases likely modulated by the intensity of transmission to dogs and by the influence on sandfly abundance (Ben-Ahmed et al., 2009). Distribution of sandflies classically associated to VL was also shown to be dependent on bioclimate (Zhioua et al., 2007), likewise for the distribution of Phlebotomus papatasi, the vector of ZCL (Chelbi et al., 2009).

Knowledge on the epidemiological patterns and trends in leishmaniasis has much increased over the last 3 decades mainly because of a better identification of the Leishmania species involved in the natural cycles of the parasite. This knowledge greatly benefited of the availability of techniques used in typing isolates obtained from humans or from reservoirs and sandflies. In this respect, IEE or molecular techniques allowed a more precise identification of the Leishmania at the species, zymodeme and genomic (schizodeme) level (Ben Hammouda et al., 2000; Ben Ismail et al., 1992; Ben Said et al., 2006; Guerbourg et al., 2007; Guizani et al., 1993, 1994a, 1994b; Guizani et al., 2002; Hanafi et al., 2001; Kallel et al., 2008b).

As far as zymodemes are concerned, it was shown that L. major was much more homogeneous than L. infantum. Indeed, all L. major strains obtained from humans, rodents and sandflies proved to be identical to the L. major MON 25 reference strain (Aoun et al., 2008; Ben Ismail et al., 1986; Haouas et al., 2007; Kallel et al., 2005). In contrast, at least three L. infantum zymodemes occur in Tunisia: i) the MON 1, mainly causing VL and to a lesser extent SCL; ii) the MON 24, the most common agent of SCL, that also causes VL; iii) the MON 80, which is very rarely isolated (Aoun et al., 1999, 2008; Bel Hadj et al., 2000, 2002; Gramiccia et al., 1991; Haouas et al., 2007). These findings show that a single zymodeme may cause quite different diseases; and raise the as yet poorly documented question of pathogenesis and virulence of strains in cause. In addition, it was shown that the distribution of both MON 1 and MON 24 L. infantum zymodemes was different according to the geographical areas the patients originated from. Indeed, L. infantum MON 24 zymodeme was more frequently reported in VL cases from Kairouan governorate as compared to northern VL foci where MON 1 zymodeme is predominant. This was attributed to the emergent character of Kairouan foci which may favour and select atypical or rare variants (Aoun et al., 2008; Ben Abda et al., 2009). Involvement of different
sandfly species in transmission of different \textit{L. infantum} parasites can not be excluded (Guerbouj et al. 2007). It is well established that different sandfly species belonging to the sub-genus \textit{Larroussius} are involved in the transmission of \textit{L. infantum}; in Tunisia clear association between sandfly species and bioclimate was demonstrated and presence of species of this subgenus in the different bioclimatic zones allowed explaining the extension of VL (Zhioua et al., 2007).

The reservoir of \textit{L. killicki} is still debated. Some findings argue for its zoonotic origin as the disease is hypoenemic and frequently reported in rural populations. In addition, \textit{L. killicki} in Kenya was isolated from hyracoides (Sang et al., 1994). However, CCL has also been reported from urban areas and from regions where \textit{L. major} ZCL is highly prevalent and additional CCL cases were probably misdiagnosed. So, the CCL may be more frequent than previously thought. On the other hand, it was hypothesized that competition between \textit{L. major} and \textit{L. killicki} may lead to some degree of pressure exerted by the first species with a subsequent reduction in the incidence of the second one (Aoun et al., 2008). However, parasite identification still remains circumstantial; parasite isolation from cutaneous lesions is often contaminated by overgrowing microorganisms. In this situation, isolating and typing of much more additional strains is highly needed, in order to get further insight in the knowledge of the concerned \textit{Leishmania} cycles.

Even though IEE has been demonstrated to be quite efficient in identifying isolates and still constitutes a reference tool, molecular techniques would be more relevant in assessing variability of strains, as even minor genomic variations may have important consequences on the epidemiological and the clinical levels. An additional advantage of the molecular techniques is their rapidity in that culture of isolates is no more needed. On the other hand, they are much adapted for both diagnostic (PCR, Multiplex PCR, Real time- PCR) and epidemiological (PCR-RFLP, Multiplex PCR, RAPD) purposes (Aoun et al., 2008; Bel Hadj et al., 2002; Ben Ismail et al., 1992; Ben Said et al., 2006; Chargui et al., 2005; Guerbouj et al., 2007; Guizani et al., 1994a, 1994b; Guizani et al., 2002; Hanafi et al., 2001; Saadi et al., in preparation). In addition, molecular techniques are specific and highly sensitive, allowing identification of parasite species hard to isolate or to maintain \textit{in vitro} in culture or too scarce to be detected or cultured. It has been supposed that the dog is the reservoir of \textit{L. infantum} MON 24 zymodeme and \textit{P. perfiliewi} the phlebotomine vector. However, isolation of this zymodeme from either the dog or the sandfly has been unsuccessful, making the hypothesis questionable (Aoun et al., 2008). In this situation, molecular techniques would be much more suitable for this purpose. Using DNA tools, \textit{P. langeroni} was found infected with \textit{L. infantum} in an active transmission focus of SCL (Guerbouj et al., 2007). It may be concluded from all these findings that identification and typing of isolates are much needed in order to best assess the epidemiology and to investigate the diversity and changing patterns of leishmaniasis.

The incidence of leishmaniasis as reflected by the annually reported cases at the national level together with our own findings was shown to highly vary and fluctuate over the 25 year period study. These fluctuations were attributed first to the dynamics of the reservoirs’ populations and sandflies, as a consequence of climatic variations, environmental and ecological changes occurring in the endemic areas (see section 2); and second, to the immunization, through the repeated outbreaks, of the infected human populations, that progressively makes previously infected people in the endemic area less receptive to the infection. This was best demonstrated in the ZCL form (Ben Salah et al., 2007).
In Tunisia, human activities, agricultural development and the subsequent ecological changes mainly benefited to rodents which dramatically proliferated in areas where their populations were previously too low for the *Leishmania* cycle to be completed. This can explain that ZCL is by far the predominant and the most widespread form of leishmaniasis in Tunisia, in terms of both incidence and geographical distribution. In addition, no sustained control programme for ZCL was carried out in most endemic areas, because of the difficulties that arose in the organization and the feasibility of such programmes. This made the ZCL epidemics still going on. It is worth mentioning however that control projects may be successful, as demonstrated by the results of those launched in 1992 in Sidi Bouzid area and again in the mid 2000s in Sidi Bouzid and Sidi El Heni, that led to an obvious decrease in the ZCL incidence over the mid 1990s in Sidi Bouzid and the last few years in both regions.

As compared to SCL, the spread of VL has been more important and obvious. This finding was related, among other factors, to the ecology of the phlebotomine vectors as it was shown that *P. perniciosus* (sub-genus *Larroussius*), the vector of MON 1 *L. infantum* zymodeme is more resistant to dry climate as compared to *P. perfiliewi*, the presumed vector of MON 24 zymodeme (Aoun et al., 2008; Ben Abda et al., 2009). Given the fact that presence and distribution of the different sandfly species generally associated to *L. infantum* transmission explained extension of VL (Zhioua et al., 2007) and that the extensions seem to concern in majority MON 24 parasites (Aoun et al., 2008; Ben Abda et al., 2009; Kallel et al., 2008a), explanations may well relate to parasite features not accounted for by the mere attribution to zymodemes and/or to differential distribution of reservoirs yet to identify.

Even though molecular techniques namely PCR and its variants have been increasingly used in the diagnosis of all forms of leishmaniasis, it is to be pointed out that conventional techniques should not be neglected. In VL, bone-marrow aspirate examination was shown to be sensitive enough, even though not optimal. Examination of cytoconcentrated peripheral blood was demonstrated to be positive in nearly 50 % of VL cases and was proposed to be carried out at first in order to avoid the painful bone-marrow puncture (Ben Said et al., 1998, Chemli et al., 2006). In serodiagnosis, FAT was found to be very efficient and suitable as its sensitivity exceeded 95 % and rK39 DST, even though less sensitive, very useful because it is highly specific, easy to perform and adapted to epidemiological investigations (Saghrouni et al., 2009). PCR is much more adapted to the diagnosis of the asymptomatic and subclinical forms of the disease because of its sensitivity and the scarcity of parasites in such situation. In CL, parasitological diagnosis by demonstrating amastigotes in dermal specimens is again very adequate, even though PCR was demonstrated to be more sensitive (Chargui et al., 2005). In contrast, serodiagnosis is much less useful in CL diagnosis as antibodies’ amounts are often very low and results difficult to interpret.

One of the main objectives of this study was to assess whether the activity of our laboratory reflects the actual situation of the leishmaniasis at the country level, and whether it could constitute an indicator of public health relevance. As far as VL is considered, most patients originating from Kairouan and Sousse governorates, where are located the most active foci, were investigated for serodiagnosis in our laboratory. So, we may conclude that our findings reflect the actual incidence of VL in the region. This is further confirmed by the comparison of our results to the statistics on the annually recorded cases of VL and edited by the primary health care direction of the Tunisian Ministry of Health (Anonymous). Obviously, our findings match these reports as highest number of registered cases at the national level was in 1992 (n = 130), 1993 (n = 122) and 2005 (n = 122); and highest number of
cases diagnosed in our laboratory was as follows: 74 cases in 1992, 72 cases in 1993 and 83 cases in 2005. This is not surprising as nearly 50% or more of VL cases originate from Kairouan governorate.

Concerning CL, we have to point out that first SCL cases from central Tunisia were diagnosed in our laboratory in patients with clinical presentation found to be evocative of this form, and amastigotes, very suggestive of *L. infantum* as compared to *L. major* because of their small size (Ben Saïd et al., 2006). As mentioned in section 2, additional cases were further reported and the spread of SCL towards central and southern Tunisia confirmed. Similarly, the first CCL case originating from outside the original foci of Tataouine area was again diagnosed in our laboratory (Haouas et al., 2005); and CCL spread was further confirmed by additional reports leading to the revised geographical distribution of the disease (Bouratbine et al., 2005; Aoun et al., 2008). In our laboratory, we used to diagnose ZCL since 1986. Up to the late 1990s, the number of diagnosed cases was too moderate as compared to that of ZCL reported cases in the endemic regions (Figures 7 & 9). This, for at least, the two following reasons: i) at the beginning of the outbreak, many patients suffering from ZCL were only clinically diagnosed, so that only a few of them were addressed to the laboratory for parasitological confirmation, ii) from the 1990s, a great number of families originating from ZCL endemic areas migrated and settled in Sousse city and suburbs, where they progressively constituted a large community. Most of them used to return back to their home of origin for summer holidays where, because continuously exposed to phlebotomine bites, many of them contract leishmaniasis which is later addressed to us for diagnosis. It is worth mentioning that a high proportion of this community originates from Sidi Khelif in Ouled Haffouz delegation, Sidi Bouzid; and many of them were contaminated there and were later diagnosed in our laboratory. From 2000 and onwards, the number of ZCL patients who attended our laboratory for diagnosis dramatically increased. We consider that since this date the activity of our laboratory indirectly but adequately reflects the actual status of ZCL in central Tunisia, namely in Sidi Bouzid, Mahdia, Kairouan and Sousse governorates. This is best illustrated by our findings in 2004 where > 400 ZCL cases were diagnosed in our laboratory. In the same year, the highest incidence of ZCL (> 15000 cases) was registered at the national level, because of the three epidemic peaks that occurred in Sidi Bouzid (Sidi Bouzid city, Regueb, Menzel Bouzayène), Gafsa (Gafsa city, Sned, Mdhilla) and Kairouan (Nasrallah, Hajeb Layoun, Chhrarda) governorates (Anonymous). The peak we registered in 2001 was due to an outbreak that arose in Chahda, a small locality in Chorbène delegation (Mahdia governorate). Indeed, more than 50% of cases we diagnosed in 2001 were from this locality. A similar peak in Sidi El Heni delegation (Sousse governorate) was registered in 2006, with 107 cases. This peak was followed by a noticeable decrease as soon as control activities were carried out in the area. Similarly the decrease in ZCL incidence shown in our study over the last 5 years can be attributed to an important decrease in the transmission in Sidi Bouzid region where control activities were launched in the mid 2000s (Figure 9). This decrease was reflected on the national level over the same period (Figure 7). Last, our results confirm the seasonal occurrence of ZCL.

Our findings and results show that the activity of our laboratory reflects the situation of leishmaniasis at the regional level and at least in part at the national level, and may be an indicator of public health relevance. It may thus contribute to early alert of health authorities on changing epidemiological trends or emergence of leishmaniasis in the region. However, our activities may not reflect the actual situation and epidemiological changes occurring in farther endemic areas like Gafsa, Tozeur, Kébili, Médenine and Gabès known to include
highly active foci because very few patients from these regions seek medical facilities in Sousse hospital. Work is in progress to identify and characterize the *Leishmania* parasites actually circulating in central Tunisia using molecular DNA techniques.

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