



Spatial analysis of cutaneous leishmaniasis in an endemic area of Iran based on environmental factors

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Abstract

Leishmaniasis is a parasitic disease caused by different species of protozoan parasites. Cutaneous leishmaniasis (CL) is still a great public health problem in Iran, especially in Isfahan Province. Distribution and abundance of vectors and reservoirs of this disease is affected by different factors such as climatic, socioeconomic and cultural. This study aimed to identify the hotspot areas for CL in Isfahan and assess the relations between the climatic and topographic factors with CL incidence using spatial analysis. We collected data on the total number of CL cases, population at risk, vegetation coverage, altitude and climatic data for each district of the province from 2011 to 2015. Global Moran's Index was used to map clustering of CL cases across districts and the Getis-Ord (G_i^*) statistics was used to determine hotspots areas of the disease in Isfahan. We applied overlay analysis to assess the correlation between the climatic and topographic factors with CL incidence. We found the CL distribution significantly clustered (Moran's Index=0.17, $P<0.001$) with the Ardestan and Aran va Bidgol ($P<0.01$) districts along with the Naein and Natanz districts ($P<0.05$) to be strong hotspot areas. Overlay anal-

ysis revealed a high incidence of CL in areas with relative humidity of 27-30%, mean temperature of 15-19°C, mean precipitation of 5-20 mm, maximum wind speed about 12-16 m/s and an altitude of 600-1,800 m. Our study showed that spatial analysis is a feasible approach for identifying spatial disease pattern and detecting hotspots of this infectious disease.

Introduction

Leishmaniasis is a parasitic disease caused by several different species of protozoan parasites (Holakouie-Naieni *et al.*, 2017), and there is no definitive treatment for this disease (Nilforoushzadeh *et al.*, 2007). Two different hosts are required to complete the life cycle of the *Leishmania* parasite, where humans and some other mammals, especially canines (that also act as reservoirs), are the definitive hosts with sand flies belonging to the genus *Phlebotomus* in Old World and *Lutzomyia* in New World acting as intermediate hosts (Bates, 2007; Akhouni *et al.*, 2016; Bari and Rahman, 2016).

The least fatal form of the disease, cutaneous leishmaniasis (CL), is caused by *Leishmania major* and is the most common clinical form of disease in the Middle East. However, the disease affects an additional 60+ countries in tropical and subtropical regions (Jacobson, 2011). The global incidence is 0.7-1.2 million new cases per year worldwide (Alvar *et al.*, 2012) and around 1.7 billion people are estimated to live in areas where leishmaniasis is common (Pigott *et al.*, 2014). CL is a serious public health problem in many rural areas of Iran (Akhavan *et al.*, 2010). A total number of 589,913 cases of CL was reported between 1998 and 2013, with the annual incidence of 30.9 per 100,000 in the Iranian population (Holakouie-Naieni *et al.*, 2017). Isfahan Province in Iran is an endemic region for CL and the main vector with respect to humans is the *Phlebotomus papatasi* sand fly with the gerbil *Rhombomys opimus* being the main nonhuman reservoir (Yaghoobi-Ershadi *et al.*, 2005; Yaghoobi-Ershadi, 2012). Distribution and abundance of vectors and reservoirs of this disease is affected by different climatic, socioeconomic and cultural factors (Desjeux, 2001; Mollalo *et al.*, 2014). Moreover, unplanned urbanization and environmental changes such as irrigation, dam construction and desertification increase the risk of infection (WHO, 2008; Nilforoushzadeh *et al.*, 2014).

Recent advances in geographic information systems (GIS) and remote sensing (RS) have promoted the study of spatial epidemiology and environmental factors affecting the vector-borne diseases (Kassem *et al.*, 2012). GIS can determine resource allocation and is a valuable approach in implementation of control measures (Barbosa *et al.*, 2014). A number of epidemiological studies

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Key words: Cutaneous leishmaniasis; Geographical information systems; Climatic factor; Spatial analysis; Iran.

Acknowledgements: the authors would like to thank the CDC Department of Isfahan University and CDC, Ministry of Health and Medical Education in Tehran, Iran.

Received for publication: 21 April 2017.
Revision received: 1 August 2017.
Accepted for publication: 2 August 2017.

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Licensee PAGEPress, Italy
Geospatial Health 2017; 12:578
doi:10.4081/gh.2017.578

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around the world have used GIS-based methods in risk mapping studies and identifying endemic area of diseases (Sudhakar *et al.*, 2006; Bhunia *et al.*, 2010; Salahi-Moghaddam *et al.*, 2010; Mollallo *et al.*, 2014, 2015; Rajabi *et al.*, 2014; Hagenlocher and Castro, 2015). These techniques have been used for developing landscape predictors of sand fly abundance as an indicator of human vector contact. Bhunia *et al.* (2010) studied the effects of several topographic indexes on the prevalence of visceral leishmaniasis (VL) using GIS and RS. They investigated the relation between altitude, temperature, humidity, rainfall and the normalised difference vegetation index (NDVI) with the prevalence of the disease in the north-eastern Indian sub-continent, while Salahi-Moghaddam *et al.* (2010) studied VL in an endemic area of Iran using the GIS approach and Mollalo *et al.* (2014) focused on the relationship between the vegetation cover and occurrence of CL in Golestan Province in Iran based on satellite images.

The aims of the present study were to 1) identify the geographical distribution of CL in Isfahan Province; 2) search for hotspot areas; and 3) assess the relations between the climatic and topographic factors with CL incidence in Isfahan Province using the spatial analysis during the period 2011 to 2015.

Materials and Methods

Study area

Isfahan Province is located between 31° 43' to 34° 22' N and 49° 38' to 55° 31' E and lies in the central parts of the Iranian plateau covering an area of 107,027 km². Iran is a mountainous country mainly situated ≥1000 m above the mean sea level. The provincial capital is the historic city of Isfahan. According to the census of 2015, it consists of 23 districts with about 4,600,000 inhabitants. The province has a moderate and dry climate on the whole, and is a well-known endemic area of leishmaniasis (Figure 1).

Data collection and preparation

In order to conduct this study, information on the total number of CL cases, population at risk, climatic data, vegetation coverage and altitude were gathered for each district of the province.

Climate data

Climatic data including mean precipitation, mean temperature and mean humidity were obtained from Tehran Meteorological Center, collected from synoptic stations in Isfahan and neighbouring provinces, including Lorestan, Kohgiluyeh VA Boyerahmad, Semnan, Chaharmahal & Bakhtiari, Yazd, Fars, Qom and Markazi. These data were entered in ArcGIS, version 10.3 (ESRI Inc, Redlands, CA, USA) and dealt with in several steps: first, the yearly average of climatic data was calculated for 34 stations from April 2011 to March 2016; a point layer was created for 34 synoptic stations in a second step followed by ordinary Kriging was carried out for interpolation and calculation for all districts to derive a predicted value for unmeasured locations. Weights were based on the distance between the measured points, the prediction locations, and the overall spatial arrangement among the measured points (<http://support.esri.com>). The output for this step is a raster layer that was subsequently overlaid on the Isfahan polygon layer at the district level. Finally, the spatial analyst extension of ArcGIS, was used for calculating the average of each climatic variable. This function summarizes the values of a raster within the zones of

another dataset (zonal statistics) and reports the results as a table (ArcMap 10.3, Spatial Analyst; ESRI Inc., Redlands, CA, USA).

Vegetation coverage

Information on vegetation status of study area was obtained from the Moderate Resolution Imaging Spectroradiometer (MODIS) consisting of 16-day composites with 250-meter spatial resolution. These data were used to calculate the NDVI, one of the most commonly used measures to of landscape ecology and useful for the study of the epidemiology of vector-borne disease (Bavia *et al.*, 2005). The NDVI is related to the proportion of photo-synthetically absorbed radiation and is calculated as follows:

$$\text{NDVI} = (\text{NIR}-\text{RED})/(\text{NIR}+\text{RED}) \quad \text{Eq. 1}$$

where NIR stands for near infrared light and RED for red light (Jackson and Huete, 1991; Meijerink *et al.*, 1994). The NDVI data for Isfahan Province covering April 2011 to March 2016 was downloaded from the United States Geological Survey (USGS) website (<https://lpdaac.usgs.gov>).

Digital elevation model data

A digital elevation model (DEM) is an ordered array of numbers that represents the elevation over a specified segment of the landscape (Meijerink *et al.*, 1994). The DEM data for our study area was obtained from the Shuttle Radar Topography Mission (SRTM) through the USGS website (<https://lta.cr.usgs.gov/SRTM>) and overlaid on Isfahan polygon with district boundaries. Finally, zonal statistics was used for computing average of altitude for each district of Isfahan.

Population data

Population data for all 23 districts of Isfahan was obtained from National Bureau of Statistics of Iran.

Cutaneous leishmaniasis data

We obtained confirmed CL cases for Isfahan Province from the Isfahan University of medical sciences and Department of Communicable Disease Control (CDC) of the Iranian Ministry of Health in Tehran. The CL incidence rate was calculated as follows:

$$\text{Incidence rate} = \frac{\text{number of reported cases}}{\text{total population at risk}} \times 100,000 \quad \text{Eq. 2}$$

The total population at risk was defined as population for each district or province, which was obtained for each year.

Spatial analysis

A choropleth map was produced to show distribution of CL cases at the district level using a range of colours. Spatial data often are clustered, which means that stronger relationships may be present between proximate observations (Fotheringham and Brunston, 1999). Therefore, to explore spatial heterogeneity of disease distribution we used the Global Moran's Index in ArcGIS, version 10.3 to map the clustering of CL cases across districts in Isfahan. Moran's Index is a commonly used indicator of spatial autocorrelation (*i.e.* the correlation of a single variable between pairs of neighbouring observations) as well as non-random accumulation that indicates clustering. This index ranges from -1 to +1, where the value 1 means perfect positive spatial autocorrelation (high values

CL incidence of $0.05 < P < 0.1$ (Figure 4).

Figure 5 shows the overlay of DEM on the map of CL incidence. A high incidence of CL was found at altitudes between 600 and 1,800 m. Overlaying of the NDVI and CL incidence layers showed a high incidence of CL in areas with a low level of herb coverage (0.073-0.110) (Figure 6). We also found a high incidence of CL in areas with relative humidity of 27-30% (Figure 7), mean temperature of 15-19°C (Figure 8), mean precipitation of 5-20 mm (Figure 9) and maximum wind speed of 12-16 m/s (Figure 10).

Discussion

The present study investigated spatial patterns of CL incidence in an endemic area of Iran using the GIS and RS, during 2011-2015. The assessment of spatial characteristics of the CL cases by Moran's Index and derived Z-scores indicated that CL cases, as a whole, were clustered in the study area. We identified hotspots as well as coldspots for CL incidence, which were clustered in a specific area. The fact that the highest and lowest CL incidences were found in 2011 and 2015, respectively, could be due to interventional programmes, such as reinforced training, health education, disease surveillance and strengthened vector/reservoir control interventions, which were performed these years.

As can be seen in Figure 5, CL incidence was predominantly found at altitudes between 600 and 1,800 m, which represents moderately low-lying parts of the province and, indeed, the country as a whole, which includes large areas of 2,000 m above the sea mean level. We identified several high-risk areas of CL in middle of Isfahan, mainly in Natanz and Ardestan. Other studies have also demonstrated the high incidence of CL in Ardestan (Nilforoushzadeh *et al.*, 2014). The finding that CL was more prevalent in areas with moderately low and low altitudes is in accordance with a large number of other studies reporting an inverse relationship between altitude and CL incidence (Guernaoui *et al.*, 2006; Bhunia *et al.*, 2010; Holakouie-Naieni *et al.*, 2017). Guernaoui *et al.* (2006) collected 2,742 specimens belonging to nine phlebotomine species and reported that *P. papatasi*, the vector of *L. major*, is more common in the lowlands. Bhunia *et al.* (2010) showed the significant effect of altitude on spread and outbreaks of VL and found a high prevalence at low altitudes, while most of the highlands had only few cases. A possible explanation could be that active transmission of CL increases when the population density increases as it does with decreasing altitude (Cohen and Small, 1998). Moreover, the higher temperature, lower humidity and the character of vegetation cover in areas with lower altitude may contribute to a higher rate of CL due to the positive effect of these variables on vector populations and reservoir hosts (Salahi-Moghaddam Abdoreza *et al.*, 2010; Ali-Akbarpour *et al.*, 2012; Mollalo *et al.*, 2015).

NDVI is commonly used to separate three types of land cover: surfaces with sparse vegetation ($NDVI < 0.2$), surfaces partially covered by vegetation ($0.2 \leq NDVI \leq 0.5$) and surfaces fully covered by vegetation ($NDVI > 0.5$) (Momeni and Saradjian, 2007). In addition, it has been shown that an extensive vegetation cover provides

shade which reduces the surface temperature, while the air temperature decreases through the process of evapotranspiration (Chaithanya *et al.*, 2017); therefore, low-level vegetation coverage is almost always accompanied by higher temperatures and evaporation as well as less rainfall and lower relative humidity, a situation which provides favourable conditions for sand flies (Mollalo *et al.*, 2014; Shirzadi *et al.*, 2015). Accordingly, our study found a high incidence of CL cases in areas with low vegetation ($NDVI < 0.165$). This finding is in line with reported findings by Bavia *et al.* (2005) in Brazil, by Bhunia *et al.* (2010) in India and by Gadisa *et al.* (2015) in Ethiopia.

We also found hotspot areas in semi-arid regions with moderate levels of humidity (Figure 7). These areas have also been shown as important foci of CL in Ethiopia (Gadisa *et al.*, 2015), in India (Sudhakar *et al.*, 2006), in Brazil (Barbosa *et al.*, 2014) and several investigators in Iran have reported similar results (Ali-Akbarpour *et al.*, 2012; Shirzadi *et al.*, 2015). The contribution of humidity is, however, difficult to separate from the effect of rainfall because of the interaction of these two factors. It has been shown that peaks in rainfall can lead to reductions in sand fly numbers since excess precipitation reduces the amount of suitable resting sites for adult sand flies, limits their flight activity, and interferes with reproduction by sweeping away the eggs (Medlock *et al.*, 2014).

Annual variation between moderately high wind speed and CL distribution showed a positive relationship in our study. A recent study has shown that *P. similis*, the potential vector of *L. tropica* in Greece, is more common in areas with low (5.9–7.7 m/s) and medium (7.7–8.6 m/s) mean wind speeds compared to very low air movement (< 5.9 m/s). However, above wind speeds 8.6 m/s the risk decreased substantially ($OR = 0.8$; 95% $CI = 0.6-1.0$; $P = 0.002$) (Ntais *et al.*, 2013). An investigation in Darab District located in Fars Province in southern Iran showed that a wind speed above 3 m/s had a preventive effect on *P. papatasi*, the dominant sand fly in that area (Askari *et al.*, 2017). Contradictory results from different researches may be explained by the dual role of wind speed in disease distribution. For example, although sand fly biting opportunities is mitigated by strong winds, the flight distance can increase (Wu *et al.*, 2016). Moreover, the interaction between climatic factors in the epidemiology of vector-borne disease should be considered, since some climatic factors such as temperature, humidity and wind speed always operate together and interact with each other in nature (Seid *et al.*, 2014).

There are some limitations for this study. The CL surveillance system in Iran is a passive system, so underreporting is a strong possibility, especially in the rural areas. Furthermore, some errors may occur in the surveillance system, such as unreliable diagnosis and notification, or cases acquired in areas other than where they were diagnosed and reported. In addition, climate is only one of many groups of factors influencing vector distribution, while other factors such as vector ecology and socio-economic factors vary from one area to the other and should also be considered in the study of vector ecology. However, we assessed only climatic factors, while we fully understand that comprehensive research needs to consider also other factors, such as cultural, socioeconomic, immigration, demographic, sanitation and vector diversity.

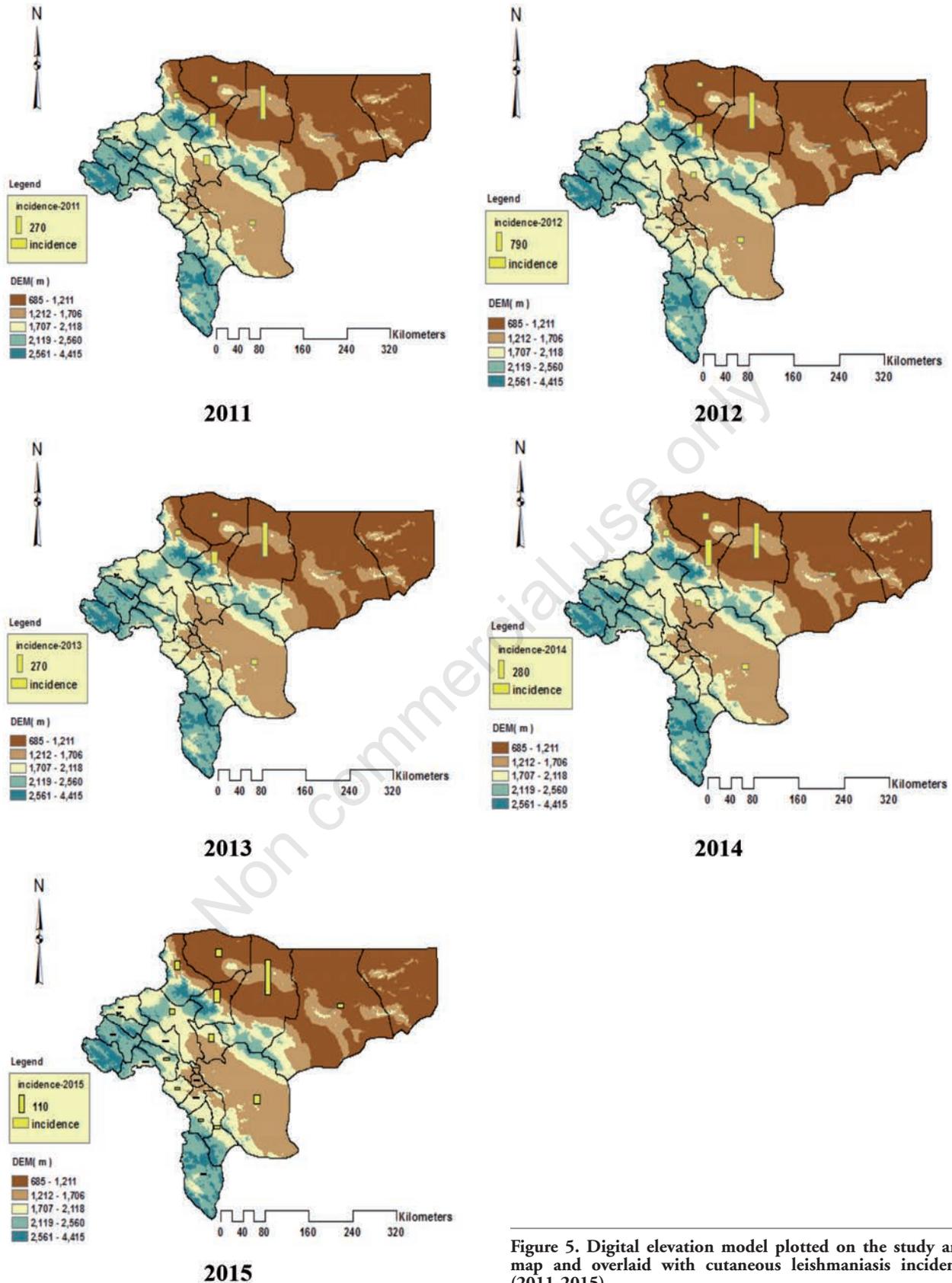


Figure 5. Digital elevation model plotted on the study area map and overlaid with cutaneous leishmaniasis incidence (2011-2015).

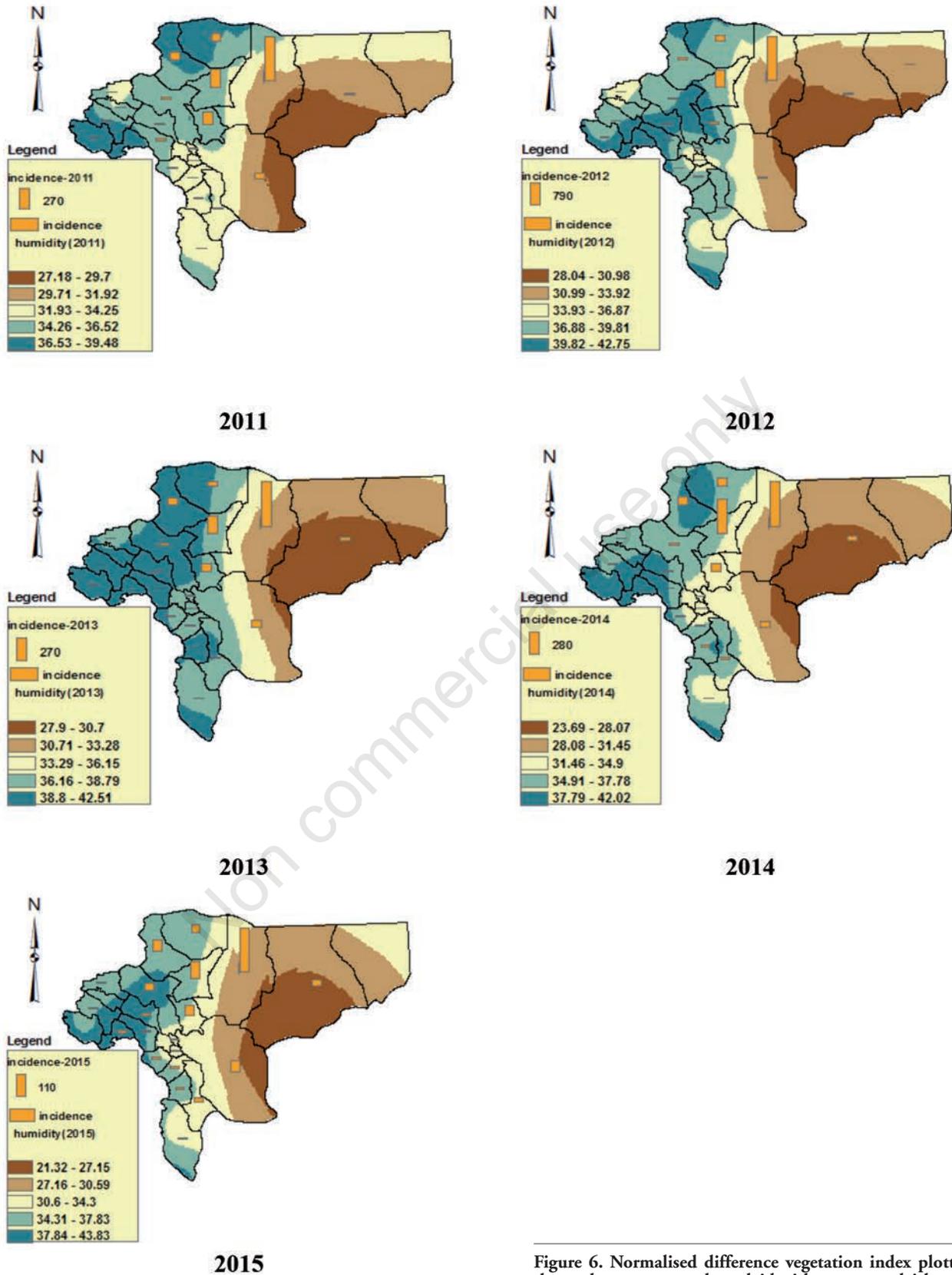


Figure 6. Normalised difference vegetation index plotted on the study area map and overlaid with cutaneous leishmaniasis incidence (2011-2015).

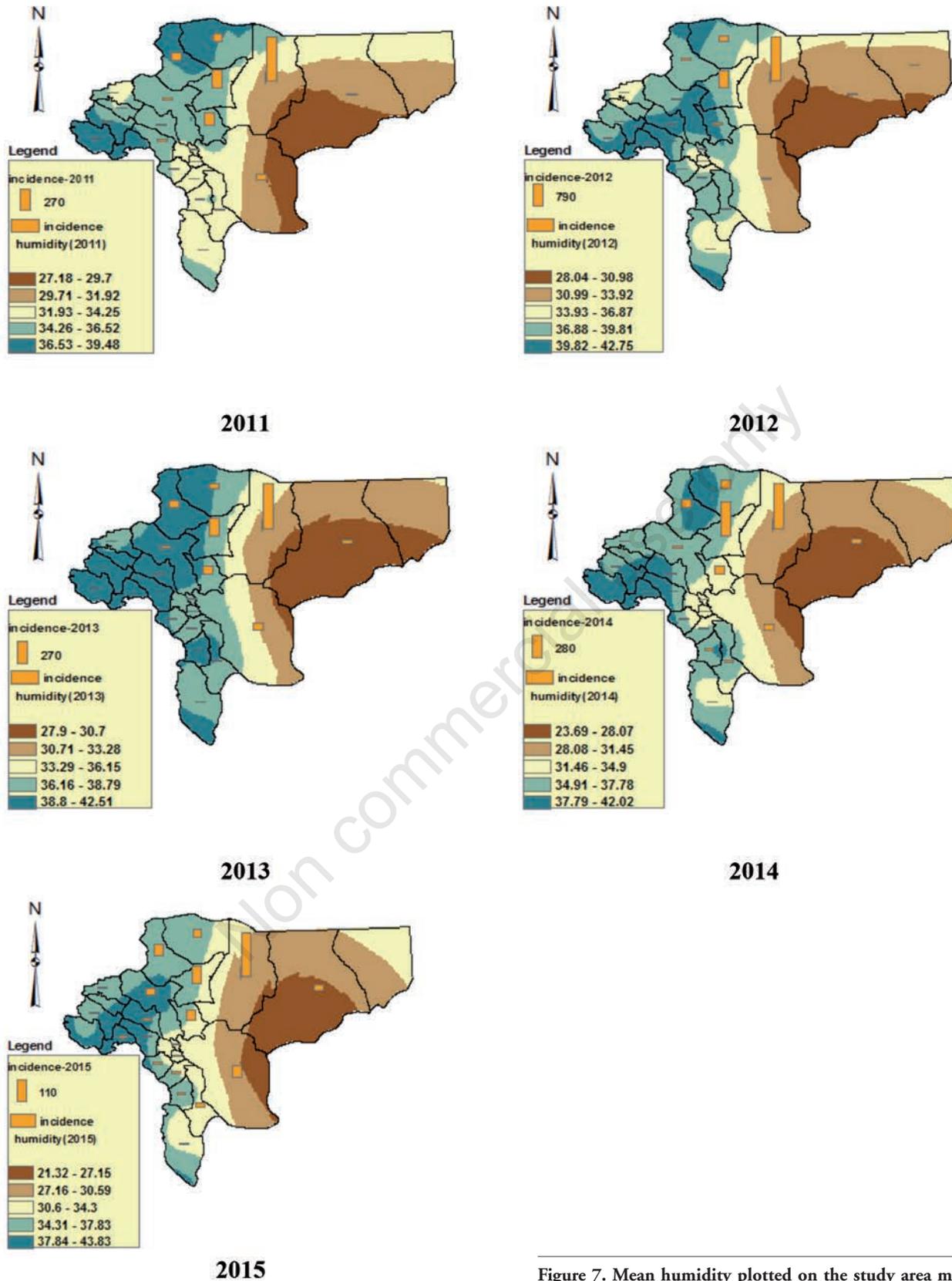


Figure 7. Mean humidity plotted on the study area map and overlaid with cutaneous leishmaniasis incidence (2011-2015).

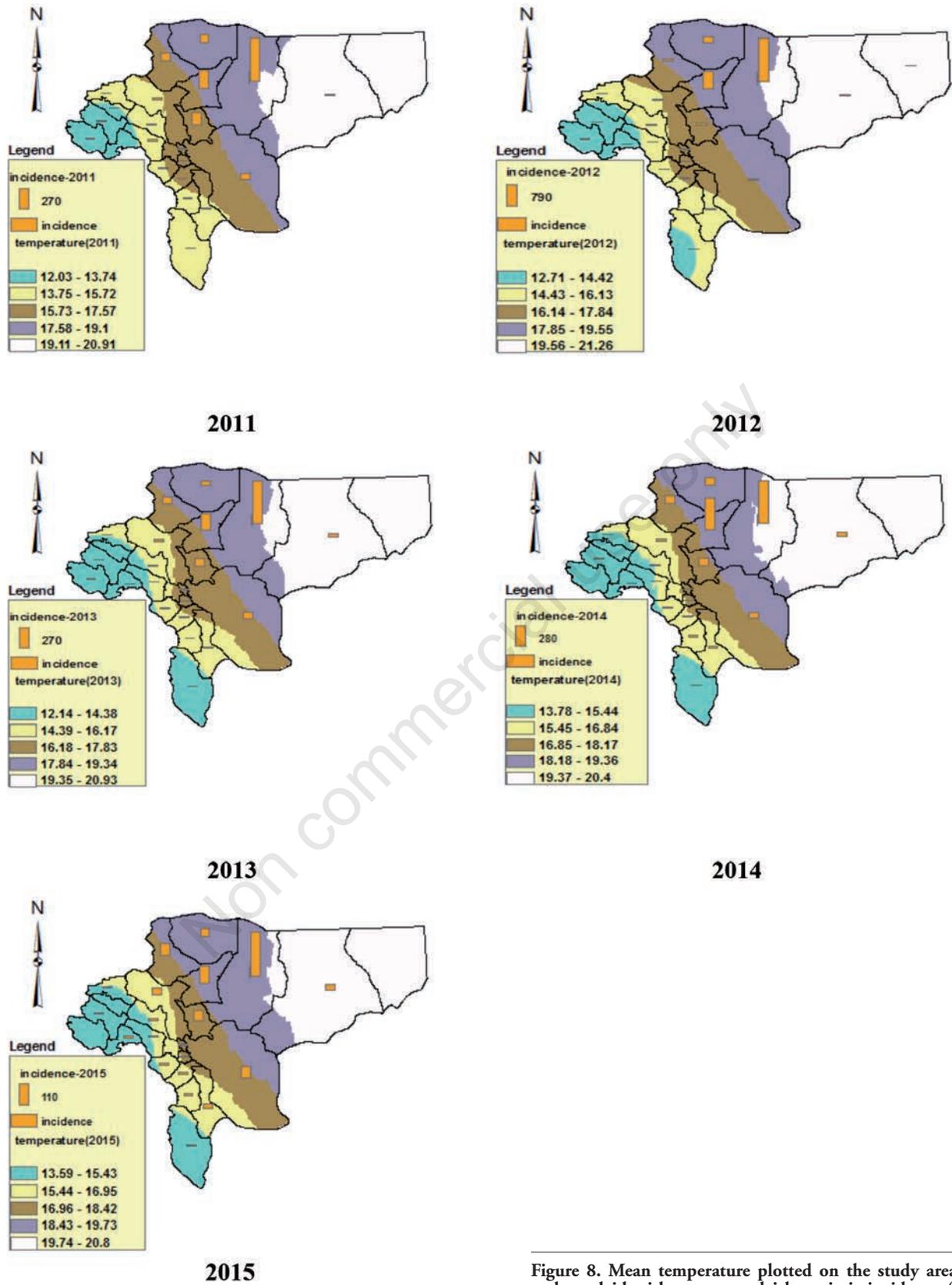


Figure 8. Mean temperature plotted on the study area map and overlaid with cutaneous leishmaniasis incidence (2011-2015).

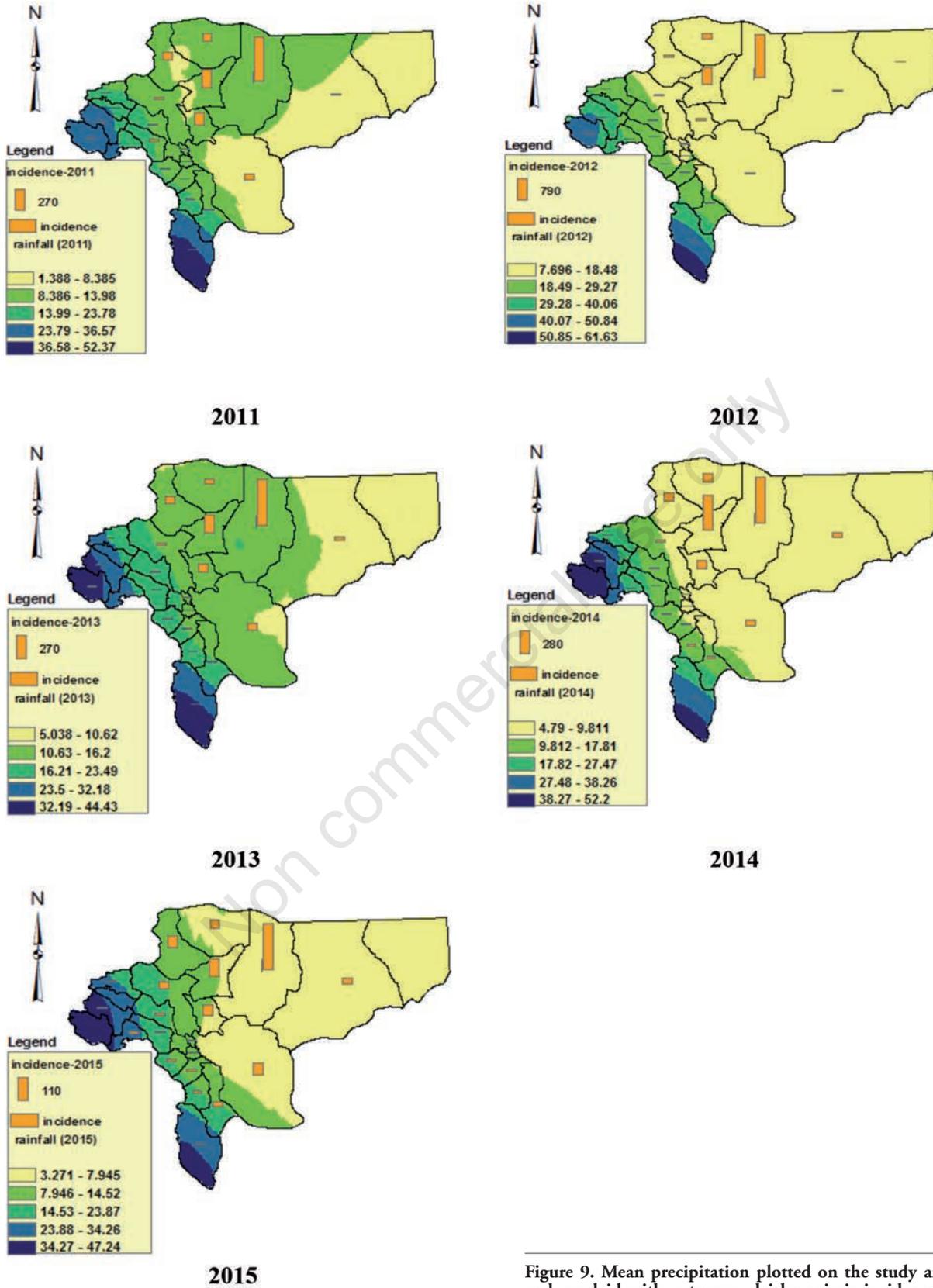


Figure 9. Mean precipitation plotted on the study area map and overlaid with cutaneous leishmaniasis incidence (2011-2015).

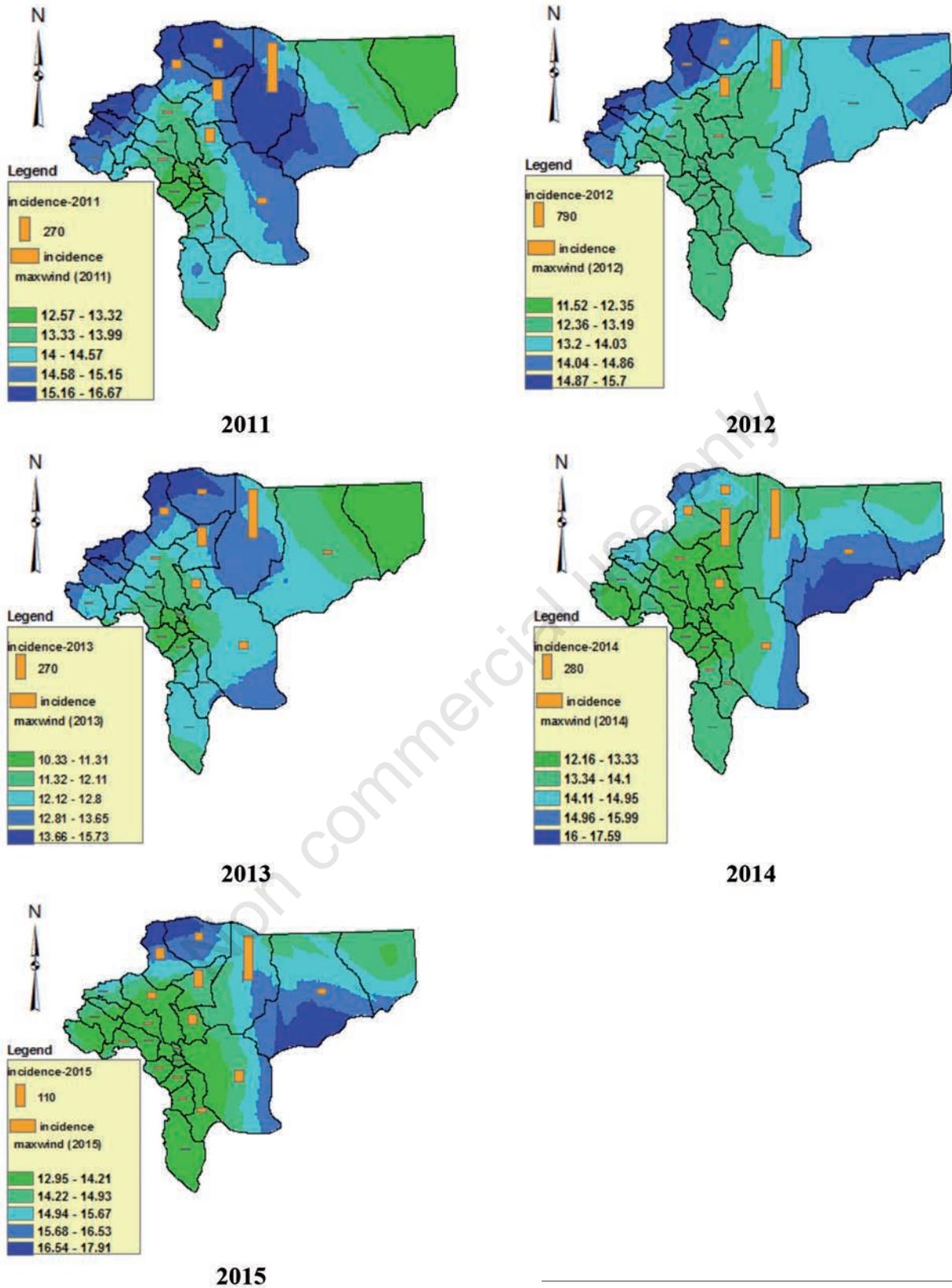


Figure 10. Mean of maximum wind speed plotted on the study area map and overlaid with cutaneous leishmaniasis incidence (2011-2015).



Conclusions

CL is a public health problem in Isfahan. Several hotspot areas were identified using spatial analysis performed by GIS and RS. Overlay analysis revealed a relationship between several climatic factors and incidence of CL in these hotspot-prone areas, the majority of which were located in semi-arid regions with low vegetation coverage. We also found fewer hotspots in lower-altitude regions with higher temperatures and less rain. In addition, a positive correlation between wind speed and hotspot areas was found. The results of the present study indicate that GIS is a feasible approach for identifying spatial disease patterns and detecting hotspots of particular infectious diseases.

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