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Abstract

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Chagas disease in European countries: the challenge of a surveillance system

L Basile¹, J M Jansà², Y Carlier³, D D Salamanca⁴, A Angheben⁵, A Bartoloni⁶, J Seixas⁷, T Van Gool⁸, C Cañavate⁹, M Flores-Chávez⁹, Y Jackson¹⁰, P L Chiodini¹¹, P Albajar-Viñas (albajarvinasp@who.int)¹², Working Group on Chagas Disease¹³

1. Department of Health, Generalitat of Catalonia, Barcelona, Spain
2. Directorate General of Public Health and Foreign Healthcare, Ministry of Health, Social Affairs and Equality, Madrid, Spain
3. Faculty of Medicine, Free University of Bruxelles, Brussels, Belgium
4. French Institute for Public Health Surveillance (Institut de Veille Sanitaire, InVS), Saint Maurice, France
5. Centre for Tropical Diseases, COHEMI network, Hospital S. Cuore, Don Calabria, Negrar, Verona, Italy
6. University Hospital Centre Careggi, COHEMI network, Firenze, Italy
7. Clinical unit of Tropical Diseases, Institute of Hygiene and Tropical Medicine, New University of Lisbon, Lisbon, Portugal
8. Department of Medical Microbiology, University of Amsterdam, Amsterdam, the Netherlands
9. National Centre of Microbiology, Instituto de Salud Carlos III, Madrid, Spain
10. Division of primary care medicine, Geneva University Hospitals and University of Geneva, Geneva, Switzerland
11. Hospital for Tropical Diseases, London, United Kingdom
12. World Health Organization (WHO) Programme on Control of Chagas disease, Department of Control of Neglected Tropical Diseases, WHO, Geneva, Switzerland
13. The members of this group are listed at the end of the article

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A study of aggregate data collected from the literature and official sources was undertaken to estimate expected and observed prevalence of *Trypanosoma cruzi* infection, annual incidence of congenital transmission and rate of underdiagnosis of Chagas disease among Latin American migrants in the nine European countries with the highest prevalence of Chagas disease. Formal and informal data sources were used to estimate the population from endemic countries resident in Europe in 2009, diagnosed cases of Chagas disease and births from mothers originating from endemic countries. By 2009, 4,290 cases had been diagnosed in Europe, compared with an estimated 68,000 to 122,000 expected cases. The expected prevalence was very high in undocumented migrants (on average 45% of total expected cases) while the observed prevalence rate was 1.3 cases per 1,000 resident migrants from endemic countries. An estimated 20 to 183 babies with congenital Chagas disease are born annually in the study countries. The annual incidence rate of congenital transmission per 1,000 pregnancies in women from endemic countries was between none and three cases. The index of underdiagnosis of *T. cruzi* infection was between 94% and 96%. Chagas disease is a public health challenge in the studied European countries. Urgent measures need to be taken to detect new cases of congenital transmission and take care of the existing cases with a focus on migrants without legal residency permit and potential difficulty accessing care.

Introduction

Chagas disease is caused by the parasite *Trypanosoma cruzi* and is considered endemic in 21 Latin American countries. It currently affects around 10 million people in Latin America, and 10 to 30 per cent of cases have developed or will develop cardiac, digestive or nervous system disorders [1]. In the last two decades many efforts have been made to reduce the incidence of Chagas disease in endemic countries [2], but exchange of population between Latin America and Europe, the United States, Australia and Japan has resulted in increased detection of *T. cruzi* in these countries [3]. In non-endemic regions, the parasite can be transmitted vertically (congenital transmission from mother to fetus), and by infected blood and organ donors [4].

In 2008, more than 38 million migrants were living in Europe, of whom 11% came from Latin America [5]. This figure did not include migrants without valid residency permit (irregular, undocumented migrants) [6], people born outside Europe who have acquired citizenship of a European country, or children from foreign countries adopted by European families. Official figures thus clearly underestimate the number of migrants from endemic areas, and therefore also the number of *T. cruzi*-infected people.

Currently, only a small number of persons infected with *T. cruzi* have been detected in Europe [4]. Several reasons account for this fact:

- Most European health professionals have little or no experience with the detection and management of Chagas disease [7].
- Access to screening programmes for the communities at risk is very limited as only a few institutions offer screening, mostly in major urban areas.
- The diagnosis of the chronic phase is usually delayed as most patients remain asymptomatic for many years [8].

There is no common European legislation to prevent the transmission of *T. cruzi* by blood donation, although in Spain and France screening of Latin American donors is mandatory, while in countries like Italy or the United Kingdom (UK) blood donation by migrants from endemic Latin American countries is prohibited and their country of origin is recorded by questionnaire [4].

Only some autonomous communities of Spain, such as Valencia [9] and Catalonia [10], have protocols for screening of pregnant women from Latin America to prevent congenital transmission. The rest of Spain and other European countries, except for some focal institutional experiences [11], have not adopted any governmental preventive measures yet.

Very few studies have estimated the prevalence of Chagas disease in European countries [12-15]. In Spain, it was estimated that between 40,000 and 65,000 residents were infected with *T. cruzi* in 2009 [4], while in other European countries the estimate range was between 12,000 and 15,000 [16].

The lack of an information system to report Chagas disease cases and transmission in all European countries makes it difficult to provide an overall figure of all diagnosed cases in Europe so far, and therefore no exact overview of the burden and public health impact of Chagas disease in Europe can be made.

For this reason, the World Health Organization (WHO) set up in 2009 a working group of experts on Chagas disease from those European countries where *T. cruzi*-positive cases had been detected (Austria, Belgium, Croatia, Denmark, France, Germany, Italy, the Netherlands, Portugal, Romania, Spain, Sweden, Switzerland and United Kingdom). The aim was to collect and assess the available information, create a network of experts to exchange information and experience between countries and define a common strategy for the epidemiological surveillance of Chagas disease [17].

This paper presents the efforts of this group of experts to provide a preliminary view of the situation in Europe, using a consensual, homogeneous methodology. The objectives of this study were to estimate the expected and observed prevalence of cases of *T. cruzi*-infected people from endemic countries of origin, the annual incidence of congenital transmission and the estimated

rate of underdiagnosis among cases of *T. cruzi* infection in 2009 in the participating countries.

Methods

Study design and population

An epidemiological study was designed to analyse aggregate measures of the prevalence of *T. cruzi* infection and the incidence of congenital transmission of Chagas disease in 2009. The units of observation were the European countries that according to the WHO estimate, had more than 400 cases of Chagas disease [4], i.e. Belgium, France, Germany, Italy, the Netherlands, Portugal, Spain, Switzerland and the UK.

Case definition

For the purposes of this study, according to the WHO case definition [18], a case of Chagas disease was considered as any individual who, as a result of a screening programme or of testing as a possible case, was positive for antibodies against *T. cruzi* in two serological (ELISA) assays.

Inclusion and exclusion criteria

The target population included three categories:

- Subjects of any age born in countries endemic for Chagas disease who were regular residents of the above-mentioned European countries in the year 2009 or the latest year for which this information was available.
- The undocumented migrants from disease endemic countries resident in the above-mentioned European countries.
- Children born in countries endemic for Chagas disease and adopted by families from the above-mentioned European countries.

Latin Americans not born in countries endemic for Chagas disease (e.g. the Caribbean islands) were excluded.

European travellers to endemic countries and cases of Chagas disease diagnosed in European travellers presumably infected in endemic countries were excluded due to the small expected number of cases and the difficulty in obtaining information about them.

Information sources

The study population was quantified using official published data obtained from national institutions in the included European countries, Eurostat and data collected by the working group and collaborators of the project. All these sources are listed in Table 1 and the corresponding data are shown in Table 2.

The numbers of diagnosed cases of Chagas disease in each European country was provided by members of the national reference institutions and members of the WHO working group.

The infection rates used to calculate the expected prevalence rate among the estimated resident population of Latin Americans in European countries (Table 3) were those published by the WHO in 2006 [19]. The rates for Bolivia were calculated according to available data on the Bolivian population living in Europe [20,21]. The rates for French Guyana and Surinam were provided respectively by the Institute of Health

Surveillance (Institut de Veille Sanitaire, France) and by the Department of Medical Microbiology of the University of Amsterdam (the Netherlands) and rely on estimations on immigrants from these countries living in Europe.

TABLE 1

Information sources for estimates of migrant residents (legal and undocumented), adoptions and annual births in nine studied European countries

Country	Category	Institution and reference year
Belgium	Legal immigration	National register, Directorate of Statistics and Economic Information (DGSIE), 2006
	Estimated undocumented immigration	Faculty of Medicine, Free University of Bruxelles, Brussels, Belgium 2006
	Adoptions	Adoption in French and Dutch-speaking Belgium, Belgian Directorate of adoption, 2001–2009
	Annual births	National register, Directorate of Statistics and Economic Information (DGSIE), 2006
France	Legal immigration	Institute of Health Surveillance (INVS), 2008
	Estimated undocumented immigration	Institute of Health Surveillance (INVS), 2008
	Adoptions	Institute of Health Surveillance (INVS), 1980–2007
	Annual births	Institute of Health Surveillance (INVS), 2008
Germany	Legal immigration	Eurostat, 2008
Italy	Legal immigration	Italian National Institute of Statistics (ISTAT), 2009
	Estimated undocumented immigration	Centre for Tropical Diseases, Hospital Ospedale Sacro Cuore Don Calabria, Verona, Italy, 2009
	Adoptions	Commission for international adoptions, Presidency of the Council of Ministers, 2000–2009
	Annual births	ISTAT, 2008
the Netherlands	Legal immigration	Statistics Netherlands, 2008
	Estimated undocumented immigration	Central government (Rijksoverheid), 2005
Portugal	Legal immigration	Statistics Portugal (INE), 2009
	Estimated undocumented immigration	Institute of Hygiene and Tropical Medicine, New University of Lisbon, Lisbon, Portugal
	Annual births	INE Portugal, 2009
Spain	Legal immigration	Statistics Spain (INE), 2009
	Estimated undocumented immigration	Statistics Spain (INE), 2009
	Adoptions	Statistics Spain (INE), 2000–2007
	Annual births	Statistics Spain (INE), 2008
Switzerland	Legal immigration	Federal department of justice and police, 2009
	Estimated undocumented immigration	Division of primary care medicine, Geneva University Hospitals and University of Geneva, Geneva, Switzerland, 2009
	Adoptions	Federal office of statistics, Section demography and migration, 1979–2008
	Annual births	Demographic portrait of Switzerland, 2008
United Kingdom	Legal immigration	Office for National Statistics, Social Surveys Dataservice, 2009
	Estimated undocumented immigration	<ol style="list-style-type: none"> 1. Sveinsson, Kjartan Páll. Bolivians In London - Challenges and Achievements of a London Community, Runnymede Community Studies, Runnymede Trust. 2007 2. Buchuck S. Crossing borders: Latin American exiles in London. Untold London, 2010 3. Bérubé M. Colombia: In the crossfire. Migration Information Source. Migration Policy Institute. 2005 4. James M. Ecuadorian identity, community and multi-cultural integration. Runnymede Trust. 2005
	Annual births	Office for National Statistics, Vital Statistics Outputs Branch, 2009

The applied rates of congenital transmission (1.4% and 7.3%) came from cohorts of migrant pregnant women living in Europe [11,22].

Data collection and analysis

To estimate the expected prevalence of *T. cruzi*-infected people in the studied countries, we first calculated the number of regular residents originating from endemic countries, according to the data published by the national statistical institutes in each country. When there were no published data, these were obtained from governmental sources or from members of the working group (Table 1).

To calculate the undocumented migrant population, we used estimates from governmental sources, national referral centres and indexed and non-indexed publications (Table 1). In the case of Spain, the official number of regular residents was subtracted from the number of migrants included in the municipal census.

In the case of children born in endemic countries and adopted by European families, we sought official data sources on adoption by country of birth (Table 1). The inclusion of this population in the study depended on the availability of data on adoptions, and finally data from five countries (Belgium, France, Italy, Spain, and Switzerland) were included.

To obtain the expected absolute number of cases of *T. cruzi* infection, the number of regular and undocumented migrants from Latin America and the number of adopted children, stratified by country of origin, was multiplied by the corresponding national infection rates in the countries of origin. A two-sided confidence intervals method with continuity correction for the single proportion [23] was applied to calculate the expected number of cases in migrants for every endemic country

of origin. The expected number of cases obtained was divided by the corresponding reference population to obtain the expected prevalence rate (shown as percentage). In the case of minimum and maximum values for reference population, an average value was applied to calculate the expected prevalence.

To calculate the observed prevalence of *T. cruzi*-infected people, the members of the working group were asked to actively search for cases diagnosed in their country up to the year 2009, dividing this amount by the total reference population to obtain the observed prevalence rate, shown as percentage.

To estimate the expected annual incidence of congenital transmission, national data on annual births of children of women from endemic areas stratified by country of birth or nationality of the mother as registered in 2009 or the latest year available was collected (Table 1). These figures were multiplied by the respective rates of infection in endemic countries, which provided an estimate of the absolute number of mothers infected with *T. cruzi* who gave birth in one year. Applying the range of congenital transmission rates (1.4% to 7.3%) to this result gave an estimate of the number of *T. cruzi*-infected children born in each participating European country. The annual incidence rate of congenital transmission in the population at risk was obtained by dividing the number of children infected in one year by the number of pregnancies in that year.

To estimate the index of underdiagnosis we calculated the rate ratio between the observed and expected prevalence rates. The result represents the proportion of diagnosed cases divided by the total estimated cases. The index is presented as a percentage obtained from the following formula: *1-rate ratio*.

TABLE 2

Estimates of migrants resident in nine studied European countries, legal and undocumented, originating from countries endemic for Chagas disease, and births to mothers from endemic countries, 2009

Country	Resident immigrants								Annual births		
	Regular population		Estimated undocumented (min–max)		Adoptions		Total (min–max)				
	Nb	%	Nb	% ^a	Nb	%	Nb	% ^a	Nb	%	
Belgium	28,880	1	14,440	1	490	1	43,810	1	722	1	
France	97,981	4	51,500	5	19,389	51	168,870	5	5,545	10	
Germany	85,313	4	Not reported	-	Not reported	-	85,313	3	Not reported	-	
Italy	260,864	12	112,000–120,000	11	6,784	18	379,648–387,648	12	3,351	6	
The Netherlands	220,172	10	17,400	2	Not reported	-	237,572	7	Not reported	-	
Portugal	110,113	5	11,011	1	Not reported	-	121,124	4	3,950	7	
Spain	1,263,342	56	484,509	47	6,354	17	1,754,205	53	35,525	67	
Switzerland	35,761	2	38,000–42,000	4	4,994	13	78,755–82,755	2	375	1	
United Kingdom	162,517	7	250,000–335,000	28	Not reported	-	412,517–497,517	14	3,433	6	
Total	2,264,943	101^b	978,860–1,075,860	99^b	38,011	100	3,281,814–3,378,814	101^b	52,901	98^b	

^a In the case of minimum and maximum values, the percentage refers to the average value.

^b The deviation is due to rounding.

Results

More than three million migrants from endemic countries (MEC) were estimated to live in the nine European countries included in the study, representing 1% of the total population living in Europe. Due to immigration from Brazil, Portugal was the country with the highest percentage of migrants coming from endemic areas. Among the countries where no Romance language is spoken, the Netherlands had the highest percentage of migrants coming from endemic countries, mainly from Surinam (84% of MEC in the Netherlands), a former Dutch colony and an endemic country for Chagas disease with a low infection rate.

Prevalence in migrants and adoptees

For details about MEC living in Europe, multiple sources of information were used (Tables 1 and 2). However, it was not possible to identify all people at risk due to the lack of data stratified by endemic country. Between 40,227 and 62,724 people infected with *T. cruzi* resided regularly in the included countries, accounting for between 1.8% and 2.8% of all regular MEC (Table 4). The highest prevalence estimation for regular MEC was seen in Spain, where between 2.3% and 3.8% of them were infected with *T. cruzi*.

The estimated numbers of undocumented MEC infected by *T. cruzi* were very high: prevalence estimations were substantially higher than for regular MEC, with the

TABLE 3

Distribution of the migrant population from countries endemic for Chagas disease resident in nine studied European countries, and estimated number of people infected, 2009

Endemic country		Infection rate	Total regular and undocumented immigrant population ^a		Estimated number of infected people ^b		
			%	Nb	% ^c	Nb	95% confidence interval
Argentina		4.13		237,678	7.1	9,815	9,626–10,006
Belize		0.74		2,464	0.1	18	11–29
Bolivia	min	10		268,926	8.4	26,893	26,597–27,188
	max	27.5		290,926		80,014	79,539–80,470
Brazil		1.02		670,299	20.1	6,837	6,703–6,971
Chile		0.99		99,483	3.0	985	925–1,045
Colombia	min	0.96		476,244	15.4	4,496	4,334–4,620
	max			546,244		5,168	5,025–5,353
Costa Rica		0.53		4,808	0.1	25	16–37
Ecuador		1.74		612,809	18.4	10,662	10,479–10,847
El Salvador		3.37		15,389	0.5	519	476–565
Guatemala		1.98		9,183	0.3	182	157–210
Guyana		1.29		23,555	0.7	13	7–24
French Guyana	min	0.25		18,987	0.6	47	36–63
	max	0.5				94	78–116
Honduras		3.05		27,121	0.8	827	773–884
Mexico		1.03		74,346	2.2	766	714–825
Nicaragua		1.14		13,317	0.4	152	129–178
Panama		0.01		4,555	0.1	0	0–5
Paraguay		2.54		87,550	2.6	2,224	2,136–2,320
Peru	min	0.69		268,957	8.2	1,856	1,775–1,936
	max			273,957		1,890	1,808–1,972
Surinam	min	0.15		183,216	5.5	287	257–330
	max	0.5				954	898–1,008
Uruguay		0.66		69,702	2.1	460	418–502
Venezuela		1.16		93,836	2.8	1,089	1,023–1,154
Undetermined ^d				19,389	0.6	165 384	
Total				3,281,814–3,378,814	100	68,318–123,078	100

^a The total immigrant population from Bolivia, Colombia and Peru is a range of values due to estimations of undocumented population.

^b Estimates based on infection rate of the country of origin.

^c In the case of minimum and maximum values, the percentage refers to the average value.

^d This number refers to adoption in France, for which no data is available stratified by endemic country, and the estimate of people infected was calculated by the Institut de Veille Sanitaire, France.

highest estimated prevalence in Spain (between 3.9% and 7.8% of undocumented MEC), and Switzerland (between 2.5% and 7.8% of undocumented MEC).

France had the highest number of positive cases among children adopted from endemic countries, although these were from countries with low infection rates. Cases represented between 0.8% and 2% of French adoptions from endemic countries. The overall expected prevalence in the participating countries ranged from 1.2% to 2.4% of total adoptions of children from endemic settings.

Congenital transmission

In the studied countries almost 53,000 children were born in 2009 from mothers originating from endemic countries. Of these, between 1,347 and 2,521 were born from mothers infected with *T. cruzi*, and there was congenital transmission in between 20 and 184 cases. This corresponds to between none and three infected children per 1,000 births to mothers at risk (Table 5). With 67% of births from mothers originating from endemic countries occurring in Spain, almost 90% all of cases of congenital transmission occurred in that country. In other countries, there were between none and six cases of congenital transmission per year.

Underdiagnosis

By 2009, 4,290 cases of infection with *T. cruzi* were diagnosed in the study countries (Table 6), and 89% of all cases were detected in Spain. The total observed prevalence rate was 0.13% of the total MEC. The lowest observed rates occurred in Germany (0.002%) and the Netherlands (0.003%) and the highest in Switzerland (0.223%).

The index of underdiagnosis shows that, in general, between 94% and 96% of expected cases were not diagnosed (Table 6). The index of underdiagnosis was lowest in Switzerland, where between 89% and 95% of expected cases were not detected, while in Germany,

the Netherlands, Portugal and the UK, more than 99% of expected cases in migrants were not diagnosed.

Overall, the Latin American nationalities with the greatest presence in Europe were Brazilians, Colombians and Ecuadorians, although most expected cases of Chagas were attributed to Bolivian migrants (Table 3).

Discussion

The Control of Chagas disease is a recent public health challenge in many countries in Europe. The reason is that it is an imported disease mainly affecting the migrated poor population from different Latin American countries who often have limited access to diagnosis and treatment of this disease. This also makes it difficult to quantify the disease impact in terms of expected cases. However, it is a challenge that requires urgent action due to the risks involved in the context of blood, organ and tissue donation, and the risk of congenital transmission to infants of infected mothers. In addition, the presence of potentially infected population groups who may present with heart, digestive tract and general disorders in the medium and long term, needs to be considered also with a view to the individual patient and the impact on clinical costs.

To quantify the European expected prevalence the authors decided to use initially the WHO official infection rates for every disease endemic country [18]. On the other hand, it was observed that all prevalence studies on Latin American immigrants living in Europe showed rates in the Bolivian community higher than the 6.75% WHO official estimated rate [20,21,24,25]. For this reason we preferred to use a more realistic range for Bolivian migrants (minimum 10.0%, maximum 27.5%) that was based on the known epidemiological situation in Europe. This choice could have introduced some bias at the methodological level by elevating the results in only one community. Nevertheless, the authors believe that this decision was necessary because the final results were closer to the reality that

TABLE 4

Estimated numbers of migrants from Chagas disease-endemic countries infected with *Trypanosoma cruzi* and expected prevalence in the nine studied European countries in 2009

Country	Legal (min-max)		Estimated undocumented (min-max)		Adoptions (min-max)		Total (min-max)	
	Prevalence		Prevalence		Prevalence		Prevalence	
Belgium	451-601	1.6-2.1	226-301	1.6-2.1	6-19	1.2-3.9	683-921	1,6-2.1
France	1,253-1,542	1.3-1.6	730-897	1.4-1.7	165-384	0.8-2	2,148-2,823	1.3-1.7
Germany	1,123-1,481	1.3-1.7	Not reported	-	Not reported	-	1,123-1,481	1.3-1.7
Italy	4,133-5,322	1.6-2	2,220-6,520	1.9-5.6	111-194	1.6-2.9	6,464-12,036	1.7-3.1
Netherlands	776-1,528	0.3-0.7	191-245	1.1-1.4	Not reported	-	967-1773	0.4-0.7
Portugal	1,141	1	114	1	Not reported	-	1,255	1
Spain	28,974-48,510	2.3-3.8	18,884-37,874	3.9-7.8	126-234	2-3.7	47,984-86,618	2.7-4.9
Switzerland	535-750	1.5-2.1	982-3,132	2.5-7.8	66-88	1.3-1.8	1,584-3971	2-4.8
United Kingdom	1,841-1,849	1.1	4,270-10,352	1.5-3.5	-	-	6,111-12,201	1.3-2.4
Total	40,227-62,724	1.8-2.8	27,617-59,435	2.7-5.8	474-919	1.2-2.4	68,318-123,078	2-3.6

professionals involved in the detection of cases see every day in health systems.

Another relevant point is that other applied national infection rates, based on the population in disease-endemic countries, do not take into account the effects of heterogeneity of the immigrant population living in Europe (i.e. age groups, socio-economic differences, rural-urban origin, etc.) and these differences are not reflected in the results.

The results of this study highlight the difficulty in obtaining accurate data on the population at risk and specific information on diagnosed cases, the lack of official national data, the underestimation of migrants in the official figures, and the lack of a system for reporting detected cases in non-endemic countries.

According to the estimations of expected cases in the different non-endemic countries, and to offer a better view of the situation, we classified the countries in

three groups. The first category includes only Spain, which accounts for almost 75% of expected cases. The second group is represented by France, Italy and the UK, while the third group is represented by the other non-endemic countries (Belgium, Germany, the Netherlands, Portugal and Switzerland). The key role played by Spain in the prevention and control of Chagas disease in Europe is not only due to the high expected prevalence of *T. cruzi* infection, but also relates to its pivotal position in the migrant flow to Europe and the cultural and linguistic proximity to Latin American countries. France has played a key role in the development of recent studies and specific interventions and regulations for Chagas disease [26], although the country had a low expected number of cases. This and the existence of French national territory in the endemic region of Latin America (French Guyana) places France in a distinctive position in the prevention and control plans for Chagas disease in non-endemic European countries.

TABLE 5

Estimated congenital transmission and prevalence rate per 1,000 pregnancies in women from Chagas disease-endemic areas, residing in nine studied European countries, 2009

Country	Annual births	Infected pregnant women (min–max)		Infected infants (min–max)	
		Number of cases	Cases per 1,000 pregnancies	Number of cases	Cases per 1,000 pregnancies
Belgium	722	10–13	14–18	0–1	<1
France	5,545	53–74	10–13	1–5	<1
Germany	Not reported	Not applicable	-	Not applicable	-
Italy	3,351	55–76	16–23	1–6	1
The Netherlands	Not reported	Not applicable	-	Not applicable	-
Portugal	3,950	40	10	1–3	<1
Spain	35,525	1,125–2,226	32–63	16–162	0–5
Switzerland	375	6–8	16–21	0–1	1
United Kingdom	3,433	58–84	17–24	1–6	1
Total	52,901	1,347–2,521	25–48	20–184	0–3

TABLE 6

Diagnosed cases, observed and expected prevalence rates and percentage of underdiagnosis of Chagas disease in migrants from endemic areas residing in nine studied European countries, up to 2009

Country	Cases diagnosed	Observed prevalence rate (%)	Expected prevalence rate (min–max, %)	Index of underdiagnosis (min–max, %)
Belgium	19	0.043	1.6–2.1	97.2–97.9
France	111	0.066	1.3–1.7	94.8–96.1
Germany	2	0.002	1.3–1.7	99.8–99.9
Italy	114	0.03	1.7–3.1	98.3–99.0
The Netherlands	7	0.003	0.4–0.7	99.3–99.6
Portugal	8	0.007	1	99.4
Spain	3,821	0.218	2.7–4.9	92.0–95.6
Switzerland	180	0.223	2–4.8	89.2–95.2
United Kingdom	28	0.006	1.3–2.4	99.6–99.7
Total	4,290	0.13	2–3.6	93.9–96.4

The observed prevalence was extremely low, compared with the expected rates, in Belgium, the Netherlands, Portugal and the UK, suggesting a lack of awareness and interventions (protocols, studies, etc) against Chagas disease in those countries. The UK, especially London where most Latin American immigrants to the UK reside [27], ranks second in Europe in terms of residents estimated to be infected with *T. cruzi* and cases of congenital transmission, with numbers nearly identical to those of Italy. These results are entirely novel and in contrast to UK estimates published in previous studies [16]. This discrepancy could be due to potential underestimation in official statistics of the Latin American population actually resident in the UK.

The study highlights the presence of positive cases in undocumented migrants, especially in Spain, Italy and Switzerland. These countries have large Bolivian communities not represented in official statistics [24,28] that makes it even harder for the national authorities to identify the population at risk. On the other hand these results can offer only an incomplete picture of the reality due to the limitations of estimating the reference population. Nevertheless the present study offers new information not included in previous studies that only included documented migrants [3,15]. The fact that being an undocumented migrant could be associated with originating from poor endemic areas with higher prevalence rates highlights the value of developing demographic studies that can contribute to providing more reliable estimates of this population.

The estimated results on underdiagnosis are a good indicator of the limited epidemiological impact of Chagas disease in the context of European health and surveillance systems. Epidemiological silence, understood as the lack of detected cases, which is common in some European countries, shows the need for greater involvement of European health authorities in controlling neglected tropical diseases, among others Chagas disease. The priority could be the implementation of screening programmes of target populations and the training of professionals in the detection of possible cases. The legislation or protocols already implemented in countries such as Spain or France would be very useful to reduce the differences in preparedness and available programmes between European countries. Such collaboration would be of help in developing a European surveillance system, which is essential for further progress in controlling Chagas disease.

The control of congenital transmission is undoubtedly one of the most important measures for the prevention and control of Chagas disease that should be addressed by surveillance systems because of the effectiveness of treatment in infants. Likewise, the establishment of regulations for blood and organ donation is essential to limit the impact of Chagas disease in countries where there is no vector transmission. Systematic screening of the risk population, at present only carried out in some regions of France, Spain and

Switzerland, should also be introduced after carrying out cost-effectiveness analyses to decide which measures could be most appropriate.

In terms of public health, the authors believe that the main proposals and challenges for European countries where cases have already been identified or that have residents from endemic areas are:

- To create an international information and surveillance system for the reporting of cases, control of transmission, exchange of information between European countries, and training of primary healthcare workers.
- To carry out studies to define the risk of congenital transmission in pregnant women from Latin America and to evaluate the impact of potential screening protocols for the control of congenital transmission according to the results obtained.
- To carry out epidemiological studies allowing for reliable estimation of true prevalence rates among immigrants resident in Europe.
- To consider systematic screening (by questionnaire or serological tests) blood, organ and tissue donors from endemic Latin American regions.
- To publish official statistics of migrants from Chagas-endemic countries in each European country containing data by regular and irregular status according to their country of origin.
- To facilitate access to diagnosis and treatment to groups of migrants at risk of being excluded from the national health systems such as undocumented immigrants.
- To reinforce the teaching on international health and tropical diseases in the curricula of health sciences in European Universities.

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Members of the Working Group on Chagas Disease in Europe:

Apart from the authors of this article, the present working group is composed of:

Dr. Mariella Anselmi (Centro per le Malattie Tropicali, Ospedale S. Cuore, Don Calabria, Negrar, Verona, Italy. COHEMI network), Dr. Luigi Gradoni (Dipartimento di Malattie Infettive, Parassitarie e Immunomediate, Istituto Superiore di Sanità, Roma, Italy), Dr. José Antonio Pérez Molina (Hospital Ramon y Cajal, Madrid, Spain), Dr. Joaquim Gascón (Hospital Clínic, Barcelona, Spain. COHEMI network), Dr. Jordi Gomez i Prat (Centre de Salut Internacional i Medicina Tropical Drassanes, Barcelona, Spain), Dr. Luisa Sánchez Serrano (Instituto de Salud Carlos III, Madrid, Spain), Dr. Magdalena García (Consorcio Hospital General Universitario de Valencia, Valencia, Spain), Dr. Bartolomé Carrilero Fernández (Hospital Universitario Virgen de la Arrixaca, Murcia, Spain), Dr. Jane Jones (Health Protection Agency, London, UK), Dr. Emmanuel Bottieau (Institute of Tropical Medicine, Antwerp, Belgium), Dr. Xavier Lescure (Hôpital Tenon, Paris, France), Dr. Pierre Ambroise-Thomas (Académie nationale de Médecine, Meylan, France), Dr. Jean Delmont (Centre de formation et recherche en médecine et santé tropicales, Marseille, France), Dr. Guillaume Le Loup (Service de Maladies infectieuses tropicales, Hôpital Tenon, Paris, France), Dr. Luc Paris (Service de Parasitologie-Mycologie, Groupe Hospitalier Pitié-Salpêtrière, Paris, France), Dr. Françoise Gay-Andrieu (Laboratoire de Parasitologie-Mycologie, Institut de Biologie - Hôtel Dieu, Nantes, France), Dr. Aldert Bart (Parasitology Section, Dept Med Microbiol, Academic Medical Center, Amsterdam, the Netherlands), Dr. August Stich (Tropenmedizin Missionsärztliche Klinik, Würzburg, Germany), Dr. Israel Molina (Hospital Vall d'Hebron, Barcelona, Spain), Dr. Carmen Muñoz (Hospital Sant Pau, Barcelona, Spain), Dr. Carmen Cabellos (Hospital de Bellvitge, Barcelona, Spain), Dr. Lluís Valerio (Unitat de Salut International Metropolitana Nord, Barcelona, Spain), Dr. Angel Lluís Ballesteros (Hospital de Badalona, Barcelona, Spain), Dr. Cristina Soler (Hospital de Santa Caterina, Girona, Spain), Dr. Toni Soriano (Hospital Joan XXIII, Tarragona, Spain) and Dr. Jean Jannin (Innovative and Intensified Disease Management Unit, Department of Control of Neglected Tropical Diseases, WHO, Geneva, Switzerland)

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