

Incidence and transmission patterns of tuberculosis among indigenous populations in Brazil

Eunice Atsuko Cunha^{1/+}, Lucilaine Ferrazoli², Lee W Riley³, Paulo Cesar Basta⁴, Michael Robert Honer⁵, Rosalia Maia⁶, Izaias Pereira da Costa⁷

¹Seção de Micobacteriologia, Laboratório Central de Saúde Pública de Mato Grosso do Sul, Campo Grande, MS, Brasil

²Núcleo de Tuberculose e Micobacteriose, Instituto Adolfo Lutz, São Paulo, SP, Brasil ³Division of Infectious Diseases & Vaccinology, School of Public Health, University of California, Berkeley, CA, USA ⁴Escola Nacional de Saúde Pública-Fiocruz, Rio de Janeiro, RJ, Brasil

⁵Universidade Católica Dom Bosco, Campo Grande, MS, Brasil ⁶Programa Nacional de Controle da Tuberculose, Ministério da Saúde, Brasília, DF, Brasil ⁷Faculdade de Medicina, Universidade Federal de Mato Grosso do Sul, Campo Grande, MS, Brasil

Approximately 10% of the Brazilian indigenous population lives in the state of Mato Grosso do Sul (MS), where a large number of new cases of tuberculosis (TB) are reported. This study was conducted to assess TB occurrence, transmission and the utility of TB diagnosis based on the Ogawa-Kudoh (O-K) culture method in this remote population. The incidence of TB was estimated by a retrospective review of the surveillance data maintained by the Notifiable Diseases Surveillance System for the study region. The TB transmission pattern among indigenous people was assessed by genotyping Mycobacterium tuberculosis isolates using the IS6110 restriction fragment length polymorphism (RFLP) technique. Of the 3,093 cases identified from 1999-2001, 610 (~20%) were indigenous patients (average incidence: 377/100,000/year). The use of the O-K culture method increased the number of diagnosed cases by 34.1%. Of the genotyped isolates from 52 indigenous patients, 33 (63.5%) belonged to cluster RFLP patterns, indicating recently transmitted TB. These results demonstrate high, on-going TB transmission rates among the indigenous people of MS and indicate that new efforts are needed to disrupt these current transmissions.

Key words: tuberculosis - genotyping - microbiology - molecular epidemiology - South American indigenous people

In Brazil, the incidence and mortality from tuberculosis (TB) decreased by 20% and 30% from 1990-2010, respectively. However, TB control remains an important challenge in the country (Barreto et al. 2011, Oliveira et al. 2013).

One particular challenge is the control of TB among remote populations, such as the indigenous populations of Brazil. The high incidence and mortality resulting from TB, initially described in the mid-20th century among indigenous people (Costa 1987), has remained unchanged (Marques & Cunha 2003, Basta et al. 2006, 2010, Bóia et al. 2009, Marques et al. 2010, Croda et al. 2012, Rios et al. 2013, Sacchi et al. 2013). It is not known if the high incidence of TB represents a high rate of current transmission or reactivation of previous TB infections.

In the 1960s, in the state of Mato Grosso do Sul (MS), Cuiá Evangelical Mission built a new hospital [Porta da Esperança Hospital (PEH)] in the municipality of Dourados to provide healthcare service for its indigenous patients. Approximately 30 years later, the local healthcare authorities initiated a program to expand case detection and TB diagnostic tests to improve treatment

among the indigenous people. As part of this program, the Ogawa-Kudoh (O-K) method to culture sputum was introduced with support from the Central Laboratory of Public Health of Mato Grosso do Sul (LACEN-MS).

The purpose of this study was to describe TB incidence and transmission patterns among indigenous people in MS during the period ranging from 1999-2001 and to assess the improvement in TB case detection based on the O-K culture method.

MS had a population of 2,078,072 people in 2000, of which 53,900 (2.6%) belonged to indigenous groups (IBGE 2011). During the study period, the Special Indigenous Health District of Mato Grosso do Sul (DSEI-MS) provided healthcare assistance to individuals of the Guarani-Kaioiwá, Terena, Kadiwéu, Kinikinawa and Guató ethnic groups. The DSEI-MS was organized into 13 Primary Care Units covering 29 municipalities, which included over 15,000 households scattered across 75 villages.

At the time of the study, PEH conducted clinical assessment and collection of biological samples for all indigenous patients suspected to have TB. The O-K culture method was implemented in September 1999.

Reported patient data were reviewed from the Notifiable Diseases Surveillance System (SINAN) for MS and from reports available at PEH covering the period from 1999-2001.

All indigenous patients who started TB treatment and were reported by the SINAN were included in the analysis to estimate incidence. Assessment of improvements in diagnosis after the introduction of the O-K culture method was based on guidelines of the Pan-Amer-

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+ Corresponding author: euniceatsuko@uol.com.br

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ican Health Organization (OMS/OPAS 2008) and on a proposal by McCarthy et al. (2008).

At the PEH laboratory, the sputum specimens were examined by microscopy (Ziehl-Neelsen staining) and were cultured after decontamination by the swab method with 4% sodium hydroxide and inoculated in O-K egg-based medium (WHO 1998). Once seeded, cultures were sent to LACEN-MS where they were incubated at 37°C. The results (negative or positive) were reported within 60 days of incubation. A drug susceptibility test (DST) was performed using Canetti's proportion method for isoniazid, rifampicin, pyrazinamide, ethambutol and streptomycin (Canetti et al. 1963).

Only cases in which *Mycobacterium tuberculosis* was isolated from culture were included in the analysis of transmission patterns. *M. tuberculosis* isolates were genotyped according to a standardised IS6110 restriction fragment length polymorphism (RFLP) method described by van Embden et al. (1993). RFLP-IS6110 gel electrophoretic banding patterns were analysed using the BioNumerics software, version 5.0 (Applied Maths, Kortrijk, Belgium). Patterns were compared according to pair groups based on non-weighted arithmetic averages and Dice coefficients. Two or more isolates were defined as belonging to a cluster when their IS6110 electrophoretic band patterns were indistinguishable. These strains were considered to have been isolated from TB patients who acquired the infection recently. Strains with profiles in which the similarity index was equal to or higher than 65% were grouped as belonging to a family.

This study was part of a wider research project titled Epidemiological Assessment and Operationalisation to Diagnose Tuberculosis in the Guarani-Kaiowá Population in the state of Mato Grosso do Sul. This project was approved by the Research Ethical Committee of the Federal University of Mato Grosso do Sul and by the National Commission of Research Ethics (126/2001).

According to the data available from SINAN, 3,093 TB cases were reported from MS from 1999-2001, of which 610 (~20%) occurred among indigenous individuals. Of this total, 562 (92.1%), 33 (5.4%) and 15 (2.5%) were cases of pulmonary, extrapulmonary and mixed-form disease in indigenous patients vs. 2,108 (85.3%), 319 (12.9%) and 43 (1.7%) in non-indigenous patients, respectively. Most individuals (63.4%) were male. The mean age of indigenous patients was 25 years vs. 40 years among non-indigenous individuals ($p < 0.001$). The mean annual incidence, based on data for the year 2000 (median of the period), was 424.9/100,000 for the indigenous population vs. 38.1/100,000 for non-indigenous residents (48.1/100,000 residents for the state as a whole).

More than half of the cases among indigenous individuals were reported from the Primary Care Unit of Dourados, which reported an average incidence of 1,102 cases per 100,000 inhabitants. The Amambai, Antônio João and Caarapó Primary Care Units had 540, 474 and 288 cases per 100,000 inhabitants, respectively (Table I). In the study period, 797 patients were evaluated and 321 TB cases were diagnosed at PEH (52% of all indigenous notified cases). For diagnostic investigation, 1,539

TABLE I
Population, number of tuberculosis cases (all forms) and incidence rate per 100,000 residents, by Primary Care Unit of the state of Mato Grosso do Sul (MS), Brazil, 1999-2001^a

| Primary Care Unit | Population (n) | | | | Cases (n) | | | | Incidence | | | |
|-------------------|----------------|--------|--------|---------|-----------|------|------|-------|-----------|---------|-------|---------|
| | 1999 | 2000 | 2001 | Total | 1999 | 2000 | 2001 | Total | 1999 | 2000 | 2001 | Average |
| Amambai | 8,519 | 9,009 | 9,463 | 26,991 | 63 | 49 | 34 | 146 | 739.5 | 543.9 | 359.3 | 540.9 |
| Antonio João | 1,400 | 1,480 | 1,546 | 4,426 | 7 | 5 | 9 | 21 | 500.0 | 337.8 | 582.1 | 474.5 |
| Aquidauna | 6,545 | 6,711 | 6,909 | 20,165 | 3 | 3 | 2 | 8 | 45.8 | 44.7 | 28.9 | 39.7 |
| Bodoquena | 839 | 867 | 902 | 2,608 | 1 | 0 | 0 | 1 | 119.2 | 0.0 | 0.0 | 38.3 |
| Bonito | 500 | 524 | 540 | 1,564 | 2 | 0 | 0 | 2 | 400.0 | 0.0 | 0.0 | 127.9 |
| Caarapó | 4,491 | 4,742 | 4,962 | 14,195 | 16 | 16 | 9 | 41 | 356.3 | 337.4 | 181.4 | 288.8 |
| Campo Grande | 3,055 | 3,157 | 3,241 | 9,453 | 1 | 2 | 0 | 3 | 32.7 | 63.4 | 0.0 | 31.7 |
| Dourados | 9,222 | 9,741 | 10,167 | 29,130 | 97 | 136 | 88 | 321 | 1,051.8 | 1,396.2 | 865.5 | 1,102.0 |
| Iguatemi | 3,984 | 4,178 | 4,369 | 12,531 | 7 | 8 | 6 | 21 | 175.7 | 191.5 | 137.3 | 167.6 |
| Miranda | 5,796 | 5,956 | 6,134 | 17,886 | 14 | 5 | 13 | 32 | 241.5 | 83.9 | 211.9 | 178.9 |
| Paranhos | 2,890 | 3,048 | 3,203 | 9,141 | 4 | 2 | 1 | 7 | 138.4 | 65.6 | 31.2 | 76.6 |
| Sidrolândia | 2,593 | 2,672 | 2,773 | 8,038 | 2 | 1 | 0 | 3 | 77.1 | 37.4 | 0.0 | 37.3 |
| Tacuru | 2,523 | 2,654 | 2,753 | 7,930 | 5 | 2 | 6 | 13 | 198.2 | 75.4 | 217.9 | 163.9 |
| Total | 52,357 | 54,739 | 56,962 | 164,058 | 222 | 229 | 168 | 620 | 424.0 | 418.3 | 294.9 | 377.3 |

^a: data only to indigenous population who was assisted by primary health units. Source: Notifiable Diseases Surveillance System/Secretary of Health-MS and Information System of Care for Indigenous Health/National Health Foundation.

sputum smears and 1,441 cultures were performed, of which 126 samples tested positive for *Mycobacterium*. *Mycobacterium* organisms were identified in 126 specimens from 99 patients. Four of the 126 samples were *Mycobacterium* spp other than *M. tuberculosis*.

Of the 99 culture-positive cases, 43 were smear-negative and 55 were smear-positive. In one of the positive culture samples, the smear test was not performed. Of the 1,441 cultures performed, 16 (1.1%) were smear-positive and culture-negative, nine (0.6%) were smear-positive and culture-contaminated and 78 (5.4%) were contaminated. The culture detection method increased the TB diagnosis frequency by 34.1%. Of the 90 *M. tuberculosis* isolates subjected to DST, all were sensitive to the five drugs tested.

RFLP-IS6110 genotyping was performed on 52 (52.2%) *M. tuberculosis* isolates from 99 patients. The number of IS6110 copies (bands) ranged from four-17 among these isolates. Two isolates contained four bands in the same position, suggesting contamination by two strains and were excluded from the analysis (Figure).

Of the genotyped isolates, 52 were isolated from indigenous patients, while seven were obtained from non-indigenous patients. Thirty-three (63.5%) belonged to cluster profiles A, B, B1, D, E, G, G1, G4 and G5. The number of strains per each cluster ranged from two-eight. Twenty-nine strains were distributed into only three families (A, B, G), totalling nearly 56% of all analysed strains (Table II). There were no differences in the mean age or gender according to cluster vs. non-cluster RFLP patterns.

A noteworthy feature was that in the village of Bororó, four strains of the B family and five strains of the G family were identified. In the village of Jaguapiru, two group D strains were isolated and in the village of Itaum, two G1 strains were found. One group A and two group E strains were identified in the village of Amambai. These findings allowed an epidemiologic connection to be established for 16 (30.8% of 52) indigenous individuals (Table II). The number of available strains from non-indigenous patients that were genotyped (n = 7) was too small to make any conclusions about transmission patterns.

This investigation confirmed that TB is a major healthcare concern for the indigenous populations of MS and demonstrated that the incidence during the study period was far greater than the national and regional averages or the incidence among non-indigenous people (Croda et al. 2012, Oliveira et al. 2013). Based on genotyping test results, a high rate of recent transmission appears to be occurring in indigenous subjects.

The implementation of the swab method with inoculation in O-K medium increased case detection by 34.1% compared to smear microscopy alone.

The use of this method for diagnosing TB in an indigenous population in Brazil for the first time was based on its ease of use because centrifugation is not required and the risk of contamination posed to healthcare professionals is lower. Furthermore, the method ensures incubation for up to 20 days after seeding, without com-

promising performance (Sushemihl et al. 1993, Honscha et al. 2008, Jaspe et al. 2009, Rivas et al. 2010, Palaci et al. 2013).

The high level of genotype similarity among *M. tuberculosis* strains within indigenous villages in Amambai and Dourados differs from the patterns described for other locations in Brazil (Fandinho et al. 2000, Ferrazoli et al. 2000, Suffys et al. 2000, Valim et al. 2006) and other countries (Frieden et al. 1996, Ferdinand et al. 2003, Hu et al. 2011), although these studies have focused on multidrug-resistant strains.

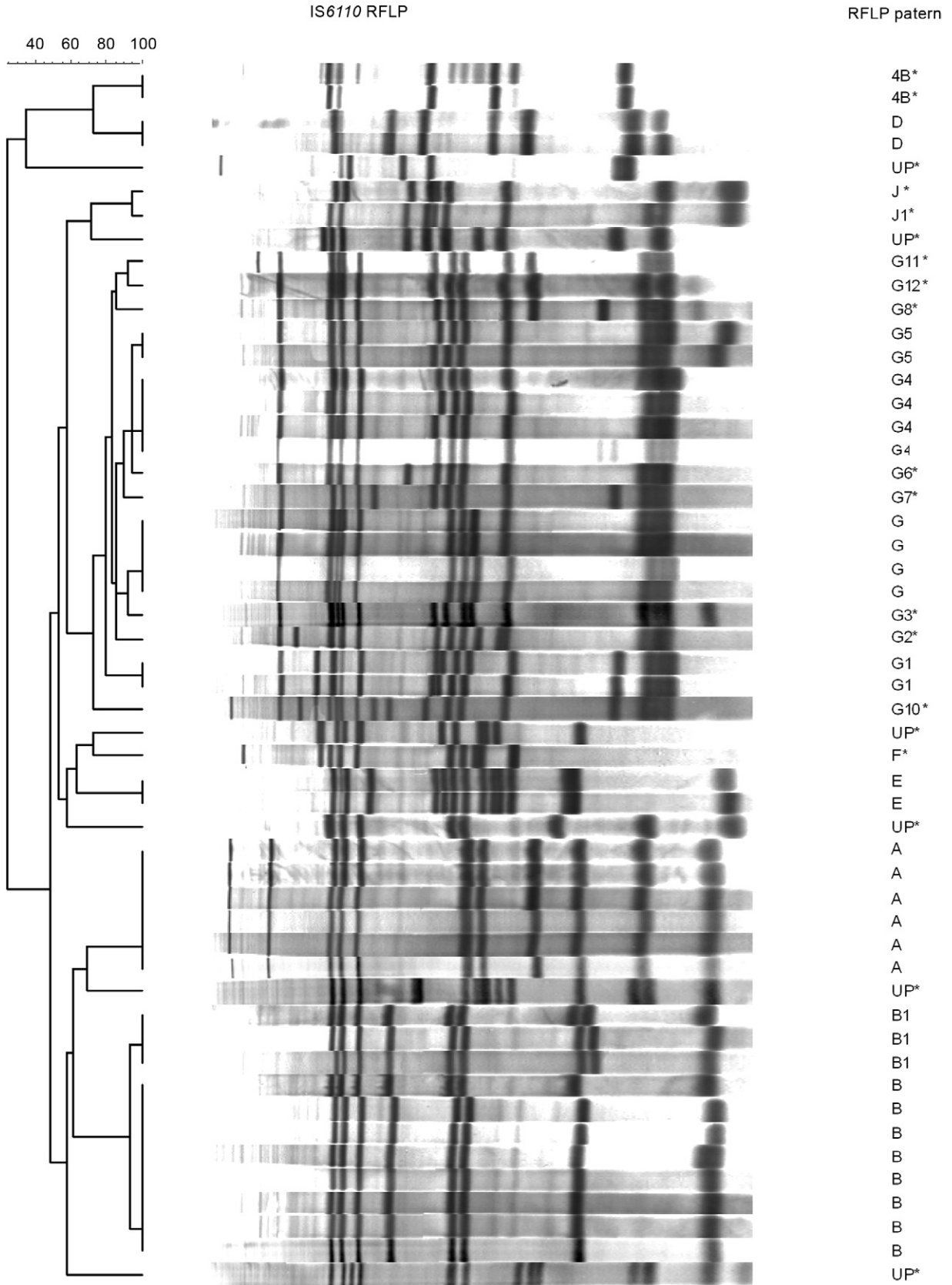
Comparable levels of similarity between strains have been reported among the Inuit in Canada (76%) (Nguyen et al. 2003) and the Waarao in Venezuela (78%) (Maes et al. 2008), in whose indigenous communities TB prevalence is higher than among the general population.

One limitation of this study is that due to inadequate storage of the isolates (frozen to -20°C), it was not possible to perform genetic analyses on all *M. tuberculosis* isolates. It is possible that other strain types circulate in the region, but that they were not identified in this study. In addition, not all reported cases used to estimate the TB incidence were confirmed by bacteriological examination.

As a result of these findings, the Dourados and Amambai Primary Care Units improved their TB control activities to interrupt TB transmission. These activities included early and high-quality diagnosis (using O-K medium), timely treatment (under dots strategy) and expanded close contact tracing. Moreover, after 2001, a preventive treatment program for latent TB infections was introduced into these communities, targeting children less than 10 years of age.

As noted by Croda et al. (2012), the above activities had a positive impact on TB control in recent years. The authors analysed not only factors predictive of failure in treatment, but also mortality in Dourados from 2002-2008. The authors concluded that there was a 90% reduction in non-completion of TB treatment among the indigenous population after dots implementation. Interestingly, the non-indigenous TB patients had 4.5 times higher mortality than the indigenous TB patients. Furthermore, Marques et al. (2010) and Santos et al. (2013) showed that the incidence of new TB cases in children less than 10 years of age decreased by approximately 40% between 2000-2006.

However, given the precarious living conditions of the indigenous populations (marked by poverty, hunger, malnutrition, anaemia, high parasite infestation and high infant mortality rates), combined with the confinement of these population to small areas (reservations, whose boundaries have been defined and approved by the federal government), TB incidence rates remain unacceptably high at approximately 200 cases per 100,000 inhabitants. Here, we have shown that the O-K culture method is simple, practical and inexpensive and that it increases case detection rates among this remote population. Nevertheless, new concerted efforts and investments are required to better characterise factors that contribute to the current high transmission frequencies among indigenous peoples from MS.



Restriction fragment length polymorphism (RFLP)-IS6110 pattern of 52 *Mycobacterium tuberculosis* isolates obtained from indigenous individuals from the state of Mato Grosso do Sul, Brazil (1999-2001). Patterns are organised by similarity according to the dendrogram to the left. Bands were aligned adopting the MT 14323 strain as the reference, using BioNumerics software, version 5.0 (Applied Maths, Kortrijk, Belgium). Asterisk means non-cluster pattern.

TABLE II

Characteristics of *Mycobacterium tuberculosis* strains by village according to cluster and non-cluster restriction fragment length polymorphism (RFLP) patterns, state of Mato Grosso do Sul, Brazil, 1999-2001

| Villages | <i>M. tuberculosis</i> isolates (n) | RFLP cluster patterns | | | | | | | | | | RFLP non-cluster patterns | | | | | | | | | | | |
|-------------|---|-----------------------|----------------|----------------|----------------|----------------|----------------|----------------|----|----|----|---------------------------|-----|-----|----|----|----|----|----|-----|---|----|-----------------|
| | | A | B | B1 | D | E | G | G1 | G4 | G5 | 4B | F | G11 | G12 | G2 | G3 | G6 | G7 | G8 | G10 | J | J1 | UP ^a |
| Amambai | 9 | 1 ^b | - | - | - | 2 ^b | - | - | 1 | - | 2 | 1 | 1 | - | - | - | - | - | - | - | 1 | - | - |
| Bororó | 13 | 1 | 3 ^b | 1 ^b | - | - | 3 ^b | - | 1 | 1 | - | - | - | 1 | - | - | 1 | - | 1 | - | - | - | |
| Caarapó | 3 | - | 1 | - | - | - | 1 | - | - | 1 | - | - | - | - | - | - | - | - | - | - | - | - | |
| Dourados | 1 | - | - | - | - | - | - | - | 1 | - | - | - | - | - | - | - | - | - | - | - | - | - | |
| Guaimbé | 2 | - | 1 | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | 1 | |
| Guassuty | 1 | - | - | - | - | - | - | - | - | - | - | - | - | - | 1 | - | - | - | - | - | - | - | |
| Itaum | 2 | - | - | - | - | - | - | 2 ^b | - | - | - | - | - | - | - | - | - | - | - | - | - | - | |
| Jaguapirú | 7 | 1 | 1 | - | 2 ^b | - | - | - | - | - | - | - | - | - | - | - | - | - | 1 | - | - | 2 | |
| Jarara | 1 | 1 | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | |
| Limão Verde | 3 | 2 ^b | - | 1 | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | |
| Miranda | 1 | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | 1 | |
| Passarinho | 1 | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | 1 | |
| Pirakuá | 1 | - | - | - | - | - | - | - | - | - | - | - | - | - | - | 1 | - | - | - | - | - | - | |
| Porto Lindo | 2 | - | 1 | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | 1 | |
| Sardinha | 1 | - | 1 | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | |
| Taquaperi | 4 | - | - | 1 | - | - | - | - | 1 | - | - | - | - | 1 | - | - | - | - | - | - | - | 1 | |
| Total | 52 | 6 | 8 | 3 | 2 | 2 | 4 | 2 | 4 | 2 | 2 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 6 |

a: unique pattern; b: individuals for whom epidemiological connection was established.

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