



Brief Report

Tick-Borne Disease Risk and Exposure among Vulnerable Populations in Perceived Non-Endemic Regions

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Abstract: Migrant and seasonal workers in the United States, among others in rural agricultural communities, may be at an elevated risk for tick-borne diseases (TBDs). This research included a survey of over 250 such workers, both women and men, in 13 locations across five eco-systems in Texas, which is generally perceived as a non-endemic state. Employing a modified Horowitz Multiple Systemic Infectious Disease Syndrome Questionnaire, a range of data was collected regarding living conditions, such as sleeping outside, along with queries about tick exposure, tick bite encounters, illnesses, and prior TBD diagnoses. Findings revealed that 67% of the respondents who scored highest on symptom severity suggestive of a TBD reported sleeping outdoors (compared to 29% of the lowest scoring), with 78% of the likely-to-highly-likely group also recalling tick-bite encounters (compared to 20% of the lowest scoring). Approximately 28% of those who reported severe symptoms also indicated previous Lyme disease diagnoses. This research serves as an initial investigation into the living conditions associated with increased risk of TBDs among vulnerable populations. It underscores the need for further assessments of TBD risk relative to vulnerable populations even in perceived non-endemic regions and highlights the paucity of actionable data as a critical public health issue.

Keywords: tick-borne disease; lyme disease; migrant and seasonal workers; vulnerable populations



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1. Introduction

Migrant and seasonal workers in agricultural and other industries may be especially vulnerable and at high-risk for contracting tick-borne diseases (TBDs), including Tickborne Relapsing Fever, Rocky Mountain Spotted Fever, and other illnesses due to occupational environments, poverty, and inferior living conditions, along with a lack of access to preventative healthcare or infectious disease physicians. Previous scholarship suggests sleeping outdoors is particularly risky for vector-borne disease exposure in general, including malaria [1]. For seasonal workers, whether in agriculture or other industries, occupational and immigration status often results in more time sleeping outdoors.

Among TBDs, Lyme disease (LD), Ehrlichiosis, and Rocky Mountain Spotted Fever particularly have spread south and west in the continental United States (US). TBD expansion is highly relevant in US states such as Texas, with increasing reports of LD cases, although officially perceived as non-endemic. Figure 1 shows US Centers for Disease Control and Prevention (CDC) reported LD cases per 100,000 in Texas, represented by color density as indicated on the right in the figure.

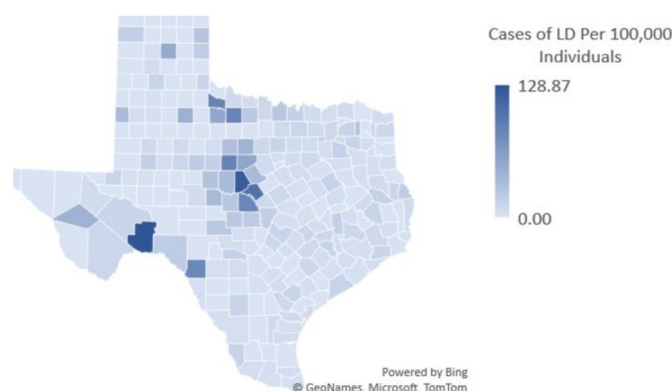


Figure 1. CDC-Reported LD Cases in Texas from 2000 to 2018 by Population Density [2].

Against this backdrop, as part of a larger ongoing effort to assess human TBD risk in different perceived endemic and non-endemic areas, we conducted an exploratory study amongst vulnerable populations living in migrant and seasonal worker communities in Texas. Survey respondents were particularly disadvantaged and vulnerable relative to occupational, immigration status, and other social factors. This brief report explores environmental, behavioral, and living conditions that may affect levels of risk among seasonal and migrant workers, among others, in agricultural communities, mostly who are Spanish-speaking and migrating from Mexico to the U.S. for employment.

2. Materials and Methods

In 2021, through a partnership with the National Center for Farmworker Health (NCFH), we conducted a survey on respondents from migrant communities, with the help of Spanish-speaking nurses and healthcare workers, across Texas. Specifically, the survey resulted in 260 validated responses. We employed the Horowitz Multisystemic Infectious Disease Syndrome Questionnaire (HMQ), recognized as a “valid, efficient, and low-cost screening tool for medical practitioners” for predicting TBDs in patients [3]. The HMQ uses the following scoring in assessing TBD likelihood: 0–20 = Not Likely, 21–36 = Possible, 36–62 = Likely, and ≥ 63 = Highly Likely [3].

Working with the NCFH, and with assistance from the Consulate General of México in Dallas in designated locations, thirteen sites were identified for survey administration (and the distribution of TBD prevention materials). The selected sites were located across nine Texas counties, covering five different ecosystems with habitats varying in suitability for relevant tick populations (Table 1):

Table 1. Survey Counties with Suitable Habitats for *Ixodes scapularis*.

Suitable for <i>Ixodes scapularis</i> Ticks	Not Suitable for <i>Ixodes scapularis</i> Ticks
<ul style="list-style-type: none"> West Gulf Coastal Plain Texas (Gregg County) Edwards Plateau (Hays, Travis counties) South Texas Brushlands (Hidalgo, Starr, Bexar, and Jim Wells counties) Oaks and Prairies Texas Ecoregion (Caldwell County) 	<ul style="list-style-type: none"> Pecos and Stalked Plains (Dallam County)

Within these ecosystems, habitat suitability is affected by varying climate and related conditions. The South Texas Brushlands (Hidalgo, Starr, Bexar, and Jim Wells counties) are notable in this regard, as survey respondents report considerable disease burden in this region, also noting an ecosystem that is particularly prone to ticks. For example, white-tailed deer are known to host ticks in the Lower Rio Grande Valley and assist in the distribution of infected ticks throughout the region [4]. Both temperature and deer density are established factors in creating suitable tick habitats [5], and deer density has been associated with the spread of Cattle Tick Fever, or Bovine Babesiosis. As shown in

Table 2, various ticks and related causative disease agents in humans and fauna are found within Texas (note: the Cayenne is limited to only the South Texas region):

Table 2. Tick Species and Diseases in Texas [6] *.

Lone Star Tick Species: <i>Amblyomma</i> <i>Americanum</i> Linnaeus	American Dog Tick Species: Dermacentor <i>Variabilis</i> (Say)	Brown Dog Tick Species: <i>Rhipicephalus</i> <i>Sanguineus</i> Latreille	Blacklegged Tick Species: <i>Ixodes</i> <i>Scapularis</i> (Say)	Gulf Coast Tick Species: <i>Amblyomma</i> <i>Maculatum</i> Koch	Cayenne Tick Species: <i>Amblyomma</i> <i>Mixtum</i> Koch	Winter Tick Species: Dermacentor <i>Albipictus</i> (Packard)
<i>Borrelia lonestari</i> —causal agent of Southern Tick Associated Rash Infection <i>Ehrlichia chaffeensis</i> and <i>E. ewingii</i> —causal agents of human ehrlichiosis. <i>Francisella tularensis</i> —causal agent of Tularemia Red meat allergy—sensitivity to “alpha-gal”, a mammalian oligosaccharide epitope Tickborne viruses— <i>Heartland virus</i> and <i>Bourbon virus</i> may be associated with this tick	<i>Rickettsia rickettsii</i> —casual agent of Rocky Mountain spotted fever <i>Anaplasma marginale</i> —casual agent of anaplasmosis in cattle <i>Theileria equi</i> —causal agent of piroplasmosis in horses, mules and donkeys	<i>Babesia canis</i> —canine babesiosis <i>Ehrlichia canis</i> —canine monocytic ehrlichiosis <i>Ehrlichia ewingii</i> —canine granulocytic ehrlichiosis <i>Rickettsia rickettsii</i> —Rocky Mountain spotted fever <i>Hepatozoan canis</i> —canine hepatoozosis	<i>Anaplasma phagocytophilum</i> —causal agent of human granulocytic anaplasmosis <i>Anaplasma marginale</i> —causal agent of anaplasmosis in cattle <i>Babesia microti</i> —causal agent of human babesiosis <i>Borrelia burgdorferi</i> —causal agent of <i>Lyme borreliosis</i> (Lyme Disease) <i>Powassan virus</i> —causal agent of Powassan encephalitis	<i>Rickettsia parkeri</i> —form of human rickettsiosis <i>Hepatozoan americanum</i> —American canine hepatoozosis <i>Ehrlichia ruminatum</i> —experimental transmission of agent of heartwater, a highly fatal African disease of ruminants known to occur in the Caribbean with <i>Amblyomma variegatum</i> , the Tropical Bont Tick.	Spotted fever group <i>Rickettsia</i> , causal agents of human rickettsiosis <i>Theileria equi</i> —causal agent of equine piroplasmosis	<i>Anaplasma marginale</i> —causal agent of anaplasmosis of cattle with transmission accomplished through trans-stadial survival of the pathogen and vector transmission either by ticks whose feeding cycle was interrupted and a second host acquired, or by trans-host migration during grooming.

* Information in quotes taken from Texas A&M University TickApp [6].

Figure 2 shows the nine survey counties compared to CDC-reported established and estimated distribution of *Ixodes scapularis* tick populations. The Texas maps are shown to provide a visual representation of survey counties and known tick distribution.

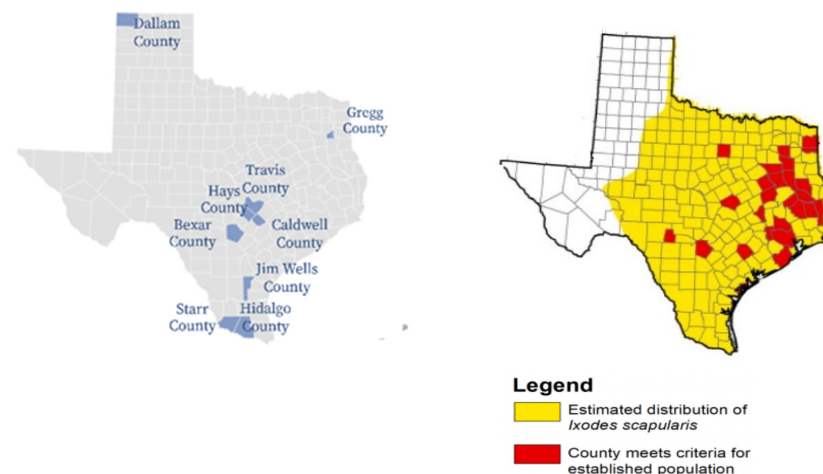


Figure 2. Survey Sites by County and *Ixodes scapularis* Tick Distribution in Texas [7].

3. Results

Of 260 total respondents included, 18 (7%) reported with scores ≥ 36 (“Likely” and “Highly Likely”); 38 (15%) scored at or between 21 and 35 (“Possibly”); and 204 (78%) scored ≤ 20 (“Unlikely” to have a TBD). Out of the 260, a total of 22% overall were possibly, likely, or highly likely to have a TBD. Living conditions and experiences, as reported, were compared among all scoring categories.

Of particular interest were survey respondents deemed likely-to-highly-likely to have a TBD based on the HMQ. Certain behaviors reported by those respondents, such as

chances of exposure to ticks through, for example, sleeping outside, may indicate higher likelihoods of disease risk. Due to similar responses and risk categories, as well as their small sample sizes compared to lower risk categories, the highly likely and likely groups were merged for the purposes of reporting descriptive results in this report.

Overall, as shown in Figure 3 and also indicated in Table 1 to highlight key findings, the Likely-to-Highly-Likely scoring respondents—and the Possibly group, to a lesser extent—were found to have higher percentages of a “previous diagnosis of LD”, “recalled a tick bite”, and “slept outside on a regular basis” with respect to their Not-Likely counterparts. In particular, tick bite recall was considerably higher among the Likely-to-Highly-Likely and Possible groups, and a previous diagnosis of Lyme disease was reported by 28% of Likely-to-Highly-Likely scoring respondents. The Likely-to-Highly-Likely group is shown in blue in Figure 3.

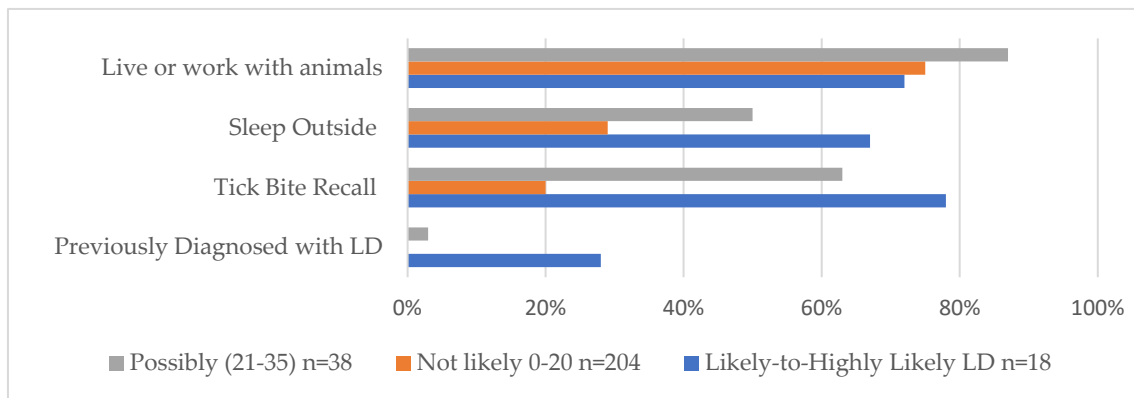


Figure 3. Survey Respondent Scores by Behavior and Tick Bite Encounter Recall.

Amongst those in each risk category reporting a tick bite encounter, reactions to the tick bite were subsequently analyzed. Respondents could select if their tick bite was associated with an Erythema Migrans rash (EM), an undefined rash, and/or was followed by flu-like symptoms, amongst numerous other symptoms. Those in the Likely-to-Highly-Likely risk category reported symptomatic and clinical presentation of a TBD at considerably higher rates than those in the Possibly or Unlikely categories, e.g., 56% percent of Likely-to-Highly Likely respondents reported a tick bite with concomitant rash and flu-like symptoms (Table 3). Almost one-third had a previous diagnosis of LD, and 67% of those most likely to have a TBD reported sleeping outdoors.

Table 3. Survey Respondent Scores by Behavior and Tick Bite Encounter Recall.

	Likely-to-Highly Likely	Possibly	Not Likely
Previously Diagnosed with LD	28%	3%	0%
Tick Bite Recall	78%	69%	20%
Tick Bite Recall—rash and flu-like	56%	14%	7.4%
Sleep Outside	67%	50%	29%
Live or work with animals	72%	87%	75%

Figure 4 shows those most likely to have a TBD (i.e., a ≥ 36 score). Out of 18 Likely-to-Highly-Likely respondents, more than half had previously received diagnoses for conditions associated to TBDs, i.e., an autoimmune disorder, Chronic Fatigue Syndrome, and/or fibromyalgia. A smaller number reported a positive LD serological test. Of the 8 respondents in this category who reported living in a LD endemic area, two were from Hidalgo County, two from Gregg County, and four from Hays County. According to CDC reports, these counties had prior LD cases: during 2000–2019, 34 confirmed cases were reported in Hidalgo, 24 in Hays, and 4 in Gregg [6].

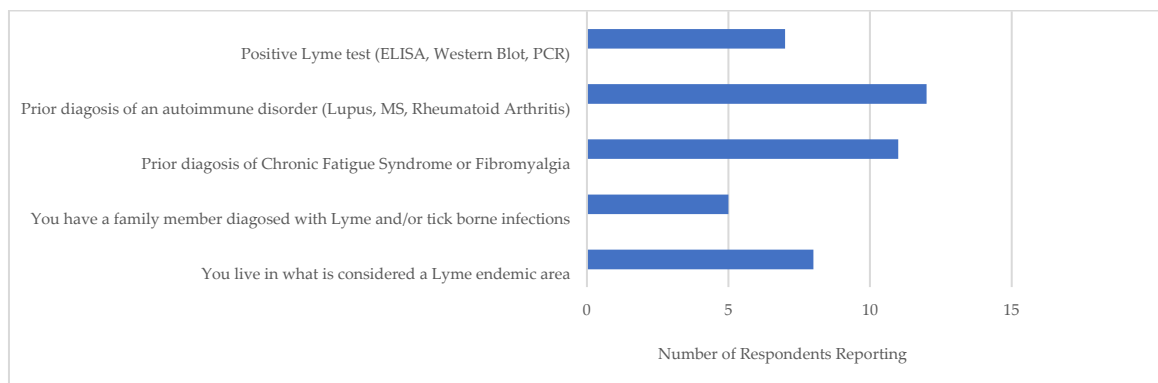


Figure 4. Frequency of Likely-to-Highly-Likely Respondent Reported TBD-Related Indicators, n = 18.

4. Discussion

This exploratory study indicates that survey respondents working in agricultural regions in Texas who scored high on the HMQ were more likely to recall a tick bite, sleep outside, and to have reported receiving a previous LD diagnosis than respondents with low scores. Living or working with animals did not appear to be associated with a higher risk of symptoms or disease in this survey. Our analysis also indicated that Likely-to-Highly-Likely respondents were found principally in the higher endemic counties amongst the survey sites. Preventative and diagnostic questions that target the main TBD symptoms, as in the HMQ survey, could assist physicians and healthcare workers in detecting and treating TBDs in these vulnerable populations, including identification of appropriate laboratories for testing. Interestingly, the highest scoring survey respondents reported previous LD diagnoses, tick bite encounters with sequela, and that they lived in TBD-endemic counties. These counties also are known for high incidences of canine TBDs such as ehrlichiosis. For example, 10.3% of all dogs tested in Hidalgo County in Texas in 2020 were positive for ehrlichiosis [8].

Vulnerable populations, especially those who may not seek treatment due to poverty, low literacy, immigration status, etc., may be at a heightened risk of TBDs in areas with reported cases. Interestingly, however, some respondents with more severe symptoms reported a prior diagnosis of LD, indicating rural and migrant clinicians may have knowledge of TBD risk in these geographic areas. Numerous TBDs with similar or overlapping symptoms are present in Texas, including Southern Tick-Associated Rash Illness (STARI), caused by the bite of an infected lone star tick, *Amblyomma Americanum*, found throughout the south-central states with distribution extending into central Texas. Along with occupational exposure, living conditions and exposure factors as reported here—such as sleeping outside or experiencing a tick bite encounter with subsequent clinical manifestations of TBDs—should be noted by migrant and rural healthcare providers and integrated into the broader epidemiological surveying infrastructure.

5. Conclusions

This research, while preliminary, suggests that vulnerable populations, in particular those whose living conditions result in increased exposure to ticks (and other disease-carrying vectors), should be considered as high risk for related diseases. Also, Lyme disease is often used as a catch-all term for TBDs and findings here suggest that the highest scoring respondents could have other tick-borne diseases that cause nonspecific and multi-system symptoms that could be mistaken for Lyme disease. The safety and health of vulnerable populations such as migrant and seasonal workers in cases of TBDs is not recognized in proportion to its scope and magnitude of risk and requires more in-depth detection and analysis at both local and broader system levels.

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C.B., C.L.M. and K.T.; writing—original draft preparation, S.P.M. and C.L.M.; writing—review and editing, S.P.M., C.L.M. and K.T.; visualization, S.P.M. and C.B.; project administration, S.P.M., C.L.M. and K.T. All authors have read and agreed to the published version of the manuscript.

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Institutional Review Board Statement: The study was conducted in accordance with the Declaration of Helsinki and approved by the Institutional Review Board (or Ethics Committee) of University of Texas at Dallas (protocol code IRB-21-149 and 8 December 2020).

Informed Consent Statement: Informed consent was obtained from all subjects involved in the study.

Data Availability Statement: Data available upon request.

Conflicts of Interest: The authors declare no conflict of interest.

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