and vinegar mixture, locally known as *kinilaw*. Other methods of fish preparation were *tinola* (boiled), *ginataan* (stewed in coconut milk), and *sinugba* (charcoal-grilled). All echinostome-infected patients had a history of having eaten snails, *kuhol* and *kiambu-ay*, prepared raw with coconut milk and lime juice (*kinilaw*), especially when found in greater abundance during the rainy season.

Human echinostome infection results from ingestion of metacercariae that encyst in secondary intermediate hosts, usually freshwater snails, tadpoles, or fish. *E. malayanum* uses various species of gastropod mollusks for primary and secondary intermediate developmental stages (2–5). Certain species of fish may also serve as secondary intermediate hosts (2). Several mollusks that may serve as primary and secondary intermediate hosts have been identified in the Philippines, including *Lymnaea* (*Bullastra*) *cumingiana*, *Radix quadraspi*, and *Physastra hungerfordiana* for *E. malayanum*, and *Pila luzonica* for *E. ilocanum* (6,7).

To our knowledge, this is the first report of *E. malayanum* infections in the southern Philippines. Local eating habits are a strong factor in echinostome infections. The general lack of awareness by health staff and the community was a big factor in the poor identification of the disease. Clinical and laboratory staff and healthcare providers need training about echinostome infections and other intestinal foodborne trematode infections. Similar environmental, sanitary, and eating practices in the region suggest that the same parasitoses should be considered to be widespread in the area. Redirecting vital resources of the local health and government units of the Caraga region to the periphery and building local capacity will help empower authorities to provide public health services in rural areas, strengthen public health programs, and further develop public health infrastructure. Moreover, a successful control program against chronic intestinal parasitoses could serve as a paradigm for local health system development of effective control measures for other endemic diseases.

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**References**


**Zoonotic Pathogens in *Ixodes scapularis*, Michigan**

To the Editor: *Ixodes scapularis*, the black-legged tick, is the predominant vector of reportable human vectorborne disease in the United States. It transmits agents that cause Lyme borreliosis, human anaplasmosis, and human babesiosis. *I. scapularis*–borne disease is becoming more frequent as this tick expands its range from tick-endemic foci in the northeastern and upper midwestern United States.

Despite Michigan’s proximity to large tick-endemic areas (Wisconsin and Minnesota to the west and Indiana to the south), active and passive surveillance data indicated that the only populations of *I. scapularis* established in the state before 2002 were in Menominee County in the Upper Peninsula (1,2). However, wildlife sampling and tick dragging in 2002–2003 suggested that *I. scapularis* had begun to invade southwestern Michigan (3), with nearby populations in northwestern Indiana (4) as the putative source.
Because we suspected these invading ticks were bringing zoonotic pathogens into southwestern Michigan, we assessed pathogen prevalence within the state’s invading and endemic I. scapularis populations. Over a 1.5-week period in April–May 2006, we collected adult I. scapularis by drag sampling at 3 recently invaded sites in southwestern Michigan and 2 tick-endemic sites in Menominee County. We targeted adult I. scapularis in the spring because this life stage has had 2 chances of becoming infected and because the adult questing peak in Michigan is greater in spring than fall (2,3).

All collected ticks were bisected aseptically, and total DNA was extracted from half after overnight lysis (DNeasy Tissue Kit; QIAGEN, Valencia, CA, USA). We used 3 PCRs to assay for Borrelia burgdorferi, B. lonestari, and B. miyamotoi (5); Anaplasma phagocytophilum (6); and Babesia spp., including Babesia microti and Babesia odocoilei (7). Borrelia-positive and Babesia-positive amplicons were purified and sequenced for species identification.

Tick densities were highest overall at tick-endemic Menominee County sites; in southwestern Michigan, they were highest at those sites closest to the putative source of the Indiana invasion. We collected 28 adult and 1 nymphal I. scapularis and 2 adult Dermacentor variabilis from tick-endemic sites. Of the adult I. scapularis, 17 (60.7%) were positive for B. burgdorferi, 4 (14.3%) were positive for A. phagocytophilum, and 2 (7.1%) were positive for Babesia odocoilei (Table). We also collected 91 adult and 10 nymphal I. scapularis and 5 adult D. variabilis from newly invaded sites. Of the adult I. scapularis, 43 (47.3%) were positive for B. burgdorferi, 1 (1.1%) was positive for A. phagocytophilum, and 4 (4.4%) were positive for Babesia odocoilei. All 4 Babesia odocoilei–positive ticks were co-infected with B. burgdorferi (this rate of co-infection was significantly greater than random expectation; p = 0.046, by Fisher exact test).

Within the tick-endemic area, comparison with prior survey data (8) indicated that the B. burgdorferi infection rate in adult ticks increased from 31.3% in 1992 to 60.7% in the present survey (p < 0.001, by Fisher exact test). A similar increasing trend was evident in the invasion area, where prevalence increased from 37.0% in 2002–2003 (at a collection site 5 km south of our southernmost site; [3]) to 47.3% in 2006. This latter trend was only marginally statistically significant due to small sample size and the short period between surveys (p = 0.046, by 1-tailed Fisher exact test).

Table. Prevalence of 3 pathogens in Ixodes scapularis ticks from 2 Michigan field sites, spring 2006

<table>
<thead>
<tr>
<th>Site</th>
<th>Life stage</th>
<th>No. ticks infected or co-infected (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>Borrelia burgdorferi</td>
</tr>
<tr>
<td>E-1</td>
<td>A</td>
<td>16</td>
</tr>
<tr>
<td>E-2</td>
<td>A</td>
<td>12</td>
</tr>
<tr>
<td>N</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>I-1</td>
<td>A</td>
<td>4</td>
</tr>
<tr>
<td>N</td>
<td>2</td>
<td>0</td>
</tr>
<tr>
<td>I-2</td>
<td>A</td>
<td>18</td>
</tr>
<tr>
<td>N</td>
<td>8</td>
<td>2</td>
</tr>
<tr>
<td>I-3</td>
<td>A</td>
<td>69</td>
</tr>
<tr>
<td>N</td>
<td>28</td>
<td>17</td>
</tr>
<tr>
<td>All endemic sites</td>
<td>N</td>
<td>1</td>
</tr>
<tr>
<td>All invaded sites</td>
<td>A</td>
<td>91</td>
</tr>
<tr>
<td>N</td>
<td>10</td>
<td>2</td>
</tr>
</tbody>
</table>

*E: endemic site; A: adult; N: nymph; I: invaded site.
†Nonsignificant level of co-infection; p = 0.378–0.640, by Fisher exact test.
‡Significant level of co-infection; p = 0.046, by Fisher exact test.
been much lower (mean 0.03 cases per 10,000 persons over same period) but appears to be increasing. Further increases in tick population size, infection, and co-infection can be expected as the invasion continues (9). Thus, medical practitioners in southwestern Michigan should be aware of the changing increasing risk for tick-borne diseases and consider disease resulting from these pathogens during diagnosis.

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