

**Ixodid Tick Effects on Deer Mice (*Peromyscus maniculatus*) Hematology and Ectoparasite  
Community Assemblages Across Populations of Varying Tick Exposure**

by

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## General Abstract

The ranges of the blacklegged tick (*Ixodes scapularis*) and the wood tick (*Dermacentor variabilis*) are expanding northward in Ontario, Canada in response to climate warming, reaching naïve or inexperienced host populations of deer mice (*Peromyscus maniculatus*) in more northern areas. As hematophagous parasites, ticks can affect their host's hematology and differing levels of hemoglobin in mice populations where tick exposure varies, may be due to the difference in exposure to ticks. If a naïve mouse population does not cohabit with an established tick population, the mice should then have higher hemoglobin levels since they are not being affected by ticks. The parasite community structures of deer mouse hosts should also differ when ticks are prevalent at varying exposure levels, as the prevalence of these ticks is expected to decrease the likelihood of other ectoparasite species co-occurring with them on the host. Blood samples were collected from individual mice from populations where: 1) both blacklegged ticks and wood ticks were prevalent, 2) only wood ticks were prevalent, or 3) where both tick species were absent in order to assess hemoglobin levels. Ectoparasites were collected from these same mice to determine parasite loads and species co-occurrences. Both my hypotheses were supported as hemoglobin levels were found to be higher in naïve mice compared to infested mice, particularly those with high tick infestations, and heterospecific parasite prevalence appeared to be higher when ticks were absent. As the ticks' ranges expand, it is important to understand the differences between naïve and experienced hosts when the prevalence of these ticks can potentially alter the physiology and community assemblages of naïve mice.

**Key Words:** Community ecology, hematology, naïve hosts, parasite-host interactions, tick range expansion

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## General Introduction

Host-parasite interactions are dynamic, as each participant influences the survival of the other. Parasites are dependent on their hosts for food and shelter and can be categorized as internal (endoparasites) or external (ectoparasites) to the host (Dougherty *et al.*, 2015). Some ectoparasites are hematophagous, whereby they negatively impact their host via blood-feeding (Bordes & Morand, 2011; Godinho *et al.*, 2013). Although this can reduce the host's fitness and can lead to host fatality, it also indirectly aids in trophic regulation and provides carrion for scavenger species (Dougherty *et al.*, 2015). When a parasite is present, the effect on the host's physiology can impact reproductive success, as less energy can be allocated to reproduction, so the number of offspring that can be produced is reduced (Fitze *et al.*, 2004). Hosts of parasites may be more susceptible to pathogenic infections (Mabbott, 2018), and may need to alter their behaviour or physiology to cope with multiple parasite infections (Thompson & Kavaliers, 1994). These energetic demands associated with parasitism can drive hosts to increase their movements, whereby the parasite manipulates the host's behaviour to favour their transmission to other potential hosts (Poulin, 2010; Thomas *et al.*, 2010; Binning *et al.*, 2017). At the same time, hosts may decrease their movements as an adaptive strategy to conserve energy (Binning *et al.*, 2014; Binning *et al.*, 2017).

*Peromyscus* mice are hosts to several ecto- and endoparasites, including ticks (Gaitan & Millien, 2016), fleas (Krasnov *et al.*, 2002), mites (Rynkiewicz *et al.*, 2013), botflies (Wolf & Baltzi, 2001), and internal nematodes (Pedersen & Greives, 2008). The parasite they host that is arguably of the greatest public health concern in North America is the blacklegged tick (*Ixodes scapularis*), which can transmit numerous zoonotic pathogens to humans (Kocan *et al.*, 2015). Despite the fact that the blacklegged tick has approximately 125 different host species that it can

feed from (Keirans *et al.*, 1996; Halsey *et al.*, 2018), *Peromyscus* mice play an essential role as a reservoir host of larval and nymph ticks (Ostfeld *et al.*, 1996). Due to their dispersal and colonizing abilities (Cummings & Vessey, 1994), *Peromyscus* mice promote the establishment of tick populations as they expand their own geographical ranges at a rate of approximately 15 km a year (Myers *et al.*, 2009). *Peromyscus* mice contract and carry the spirochaete *Borrelia burgdorferi* (the causal agent of Lyme disease) from infected blacklegged ticks and transmit the bacteria to initially uninfected blacklegged ticks when the parasites feed from them (Ostfeld *et al.*, 1996). Other tick species, such as the wood tick (*Dermacentor variabilis*) also use *Peromyscus* as a host (Minigan *et al.*, 2018). Although it has been suggested that larval ticks do not affect the survival of *Peromyscus* mice (Hersh *et al.*, 2014), physiological metrics that may be affected by tick blood-feeding, such as blood loss and iron deficiency, have not been examined for this genus.

The blood consumed by ticks contains iron-rich hemoglobin (Toh *et al.*, 2010; Pishchany & Skaar, 2012). Hemoglobin is important for oxygen transport, aiding in motor skills and immune response (Nicolas *et al.*, 2002). Little is known about the frequency of iron deficiency in wild populations of mice, although many studies have been conducted on mice in a laboratory setting with applications for human-related iron deficiency (Baggs & Miller, 1973; Nicolas *et al.*, 2002; Grant *et al.*, 2003). Baggs & Miller (1973) showed that when rats were iron deficient, they were more susceptible to bacterial infections. This pattern may also exist in natural populations that experience iron deficiency – perhaps due to the blood-feeding habits of ticks. Ticks typically feed from mice at the larval or nymph stage, and then move on to larger hosts once they mature (Estrada-Peña & de la Fuente, 2014). The cost of parasitism may be particularly high during periods that are energetically costly to mice (pregnancy, lactation), leading to insufficient

immune responses to tick infestations that cause the mice to be vulnerable to other infections (Rosales *et al.*, 1999; Dlugosz *et al.*, 2014). In a study comparing cattle (*Bos taurus*) calves infested with cattle ticks (*Rhipicephalus microplus*) and non-infested calves, there was a significant reduction in hematological parameters, including hemoglobin, leucocytes, and erythrocytes (Kaur *et al.*, 2017). Heavy parasite loads may be related to significant effects on host hematology. In several studies on avian hosts, hematocrit decreased as parasite loads increased (Møller, 1991; Whitworth & Bennet 1992; Hurtrez *et al.*, 1997). Although few experiments of this nature have been conducted on small mammals in the wild, Pfäffle *et al.* (2009) found that tick feeding on European hedgehogs (*Erinaceus europaeus*) reduced blood parameters including hemoglobin, hematocrit, and erythrocytes. Results may be similar in deer mice (*P. maniculatus*) whereby infested individuals have reduced blood parameters, and those infested with greater parasite loads have further reductions to their hematological condition.

Several studies have been conducted on how tick prevalence may affect the survival of their hosts, but physiological metrics, such as hemoglobin levels, are rarely considered. Similarly, despite many studies conducted on birds (Wheeler & Threlfall, 1986; Mallory *et al.*, 2006), mammals (Thompson *et al.*, 1998; Galloway, 2012; Webber *et al.*, 2014), and fish (Poulin, 1991; Poulin & Guegan, 2000) comparing differences in host populations where ectoparasite prevalence and intensity differ, there are few, if any, studies on mouse populations. *Peromyscus* mice are common hosts of several zoonotic pathogens that are transmitted through ticks, including *Borrelia burgdorferi*, *B. miyamotoi* (Talagrand-Reboul *et al.*, 2018), *B. bissettii* (DeNatale *et al.*, 2002), *Babesia microti* (Westblade *et al.*, 2017), and Powassan virus (Mlera *et al.*, 2018). Therefore, recognizing how *Peromyscus* hosts are physiologically affected by blood-feeding may help to make better predictions on how a host's mobility is affected by a parasite.

This provides a more accurate prediction on the ecology of the host (*i.e.*, range expansion, dispersal, mating; Binning *et al.*, 2017), which can potentially help track the transmission of pathogens they carry (Salkeld *et al.*, 2013).

Ectoparasites such as fleas (Insecta: Siphonaptera) or hematophagous mites (Acari: Mesostigmata) can co-exist with ticks (Rynkiewicz, *et al.*, 2013; Gómez-Rodríguez *et al.*, 2015). Since these ectoparasites are also blood feeders, they tend to prefer areas of exposed skin on their hosts as their micro-habitat niche (Anderson *et al.*, 2017), such as the ears (Ostfeld *et al.*, 1993; Bobbie *et al.*, 2016), which can lead to competition between species (Baer-Lehman *et al.*, 2012). Micro-habitat partitioning on a host (whereby the host is the habitat) can occur, particularly when species richness is high in an ectoparasite community and several species fit into the same niche (Gómez-Rodríguez *et al.*, 2015). However, if one species consistently outcompetes another in the micro-habitat on a host (*i.e.*, in an area both ectoparasites prefer), then prevalence or intensity differences between species may occur (Baer-Lehman *et al.*, 2012). These differences can then affect parasite species diversity on a host, which in turn can alter the effects of parasitism on the ecology and evolution of a host (Bordes & Morand, 2011).

Differences in macro-habitats or landscape ecosystems can affect the diversity of ectoparasites (Gómez-Rodríguez *et al.*, 2015). Specific geography and associated ecological factors have been shown to influence inter-individual variation in the microbiota of wild mice (Weldon *et al.*, 2015), and can potentially affect tick microbiota as well (Clow *et al.*, 2018). Latitudinal differences in parasite species richness and community structure have been studied in fish (Rohde & Heap, 1997), primates (Nunn *et al.*, 2005), and rodents (Bordes *et al.*, 2011; Preisser, 2019), where there is a trend in increasing parasite species richness with decreasing latitudes. It is also likely that there are systematic differences in genetics, habitat, and infection

status between host population sites that can affect parasite loads (Weldon *et al.*, 2015; Carbayo *et al.*, 2019). This can in turn affect interspecific competition between ectoparasite species, whereby one parasite reduces the host's susceptibility to another co-infecting parasite (Ashby & King, 2017). Ectoparasites are also able to partition their host's body, allowing for these species to co-exist across micro-habitat niches (Gómez-Rodríguez *et al.*, 2015). The influence of ticks on ectoparasite community assemblages has been widely studied and patterns of species co-occurrences tend to vary across host species depending on their habitat and ecology (Krasnov *et al.*, 2007; Krasnov *et al.*, 2009; Lareschi & Krasnov, 2010; Esser *et al.*, 2016; Krasnov *et al.*, 2020). It is important to understand how differences in macro and micro-habitats can affect parasite-host relationships when examining topics that may be affected by these parameters.

The objective of this study is to determine differences in hemoglobin levels and ectoparasite community assemblages of deer mice populations when individuals are exposed to varying established levels of two tick species; blacklegged ticks and wood ticks. Establishment levels differ by lengths of time the ticks have been established in an area, as well as the number of tick species established there. In tick established areas, deer mice have increased their grooming efficiency to counteract tick infestations (Ostfeld *et al.*, 1996; Shaw *et al.*, 2003; Hersh *et al.*, 2014), however the extent of a tick's hematological impact on *Peromyscus* hosts is still unknown. This includes differences between mice that inhabit an area where ticks have long established and less experienced mice that have not cohabited with ticks for very long – or at all. Additionally, interactions between ticks and other ectoparasites that may affect the host's hematological composition have also not been well studied. Therefore, I hypothesized that if mice are parasitized by a greater intensity of ticks, then they will have lower hemoglobin levels compared to mice infested with ticks at low intensities. Concurrently, if a naïve mouse

population has not cohabitated with an established tick population, then this mouse population should have higher hemoglobin levels relative to mice that are cohabiting with ticks since they are not afflicted by the effects of blood-feeding by these species.

The parasite community structures of the *Peromyscus* hosts may also differ when blacklegged ticks or wood ticks are prevalent, as the prevalence of these ticks are predicted to lead to a competitive relationship with other co-occurring parasites (Baer-Lehman *et al.*, 2012). This may result in a decrease in the abundance of other ectoparasite species co-occurring on the host, especially those species that congregate in the same areas on the mouse's body - *i.e.* the ears, where ticks (Ostfeld *et al.*, 1993) and mites (Bobbie *et al.*, 2016) are commonly found. I predicted that if ticks are prevalent in an ectoparasite community, then there will be fewer other ectoparasites since ticks will exclude them from the micro-habitat (the host), as they tend to prefer to feed from the same exposed areas, such as the ears. I predicted that mice parasitized by ticks would have fewer other ectoparasite species on them, as the ticks are larger and require larger blood meals (McKenzie, 1998), suggesting they are more competitive than the other common ectoparasite species found on North American small mammals that share their micro-habitat niches on the host (*i.e.* mites on ears; Bobbie *et al.*, 2016).

**Study Host Species:** *Peromyscus maniculatus* (deer mouse)

The deer mouse is one of the widest ranging mice in North America (Figure 0.1; Bedford and Hoekstra, 2015; Leo & Milien, 2017). This mouse tends to inhabit conifer and deciduous forests alike, as well as grasslands (Garcia-Elfring *et al.*, 2017). Their diets mainly consist of seeds and nuts (Cramer, 2014). Female deer mice have been documented nesting alone or in kin groups on occasion (Wolff, 1994), although parental care is exclusively maternal (Wolff, 1985).

It is during encounters with conspecifics in nests that *Peromyscus* mice typically transfer ectoparasites (such as fleas, lice, mites; Mize *et al.*, 2011; Maaz *et al.*, 2018) from one individual to another (Glicken & Schwab, 1980), however, ticks rely on one host per life stage. Ticks will emerge from the grass or forest leaf litter to make contact with mice rather than opportunistically jumping from one host to the next (Arsnoe *et al.*, 2015).

**Study Parasite Species:** *Ixodes scapularis* (blacklegged tick) and *Dermacentor variabilis* (wood tick)

Hard bodied ticks (Acari: Ixodidae) are obligate parasites that rely on vertebrate hosts for blood meals to complete the three stages in their life cycles (Figure 0.2; Brunner *et al.*, 2011; Kada *et al.*, 2017; Halsey & Miller, 2018). Several millilitres of blood are taken from the host by the tick (Brunner *et al.*, 2011), which will typically feed for three to six days as a larva or nymph (Estrada-Peña & de la Fuente, 2014). Digestion of blood starts hours after feeding, and can take several months, until moulting is complete (Estrada-Peña & de la Fuente, 2014). Since they are small and wingless, ticks rely on their hosts to move within and between habitats or to expand their geographical ranges. Therefore, ticks are constrained by the geographic barriers imposed on their hosts (Falco & Fish, 1991; Estrada-Peña & de la Fuente, 2014; Kada *et al.*, 2017).

Northward migrating birds aid blacklegged ticks in expanding their geographic ranges during the spring (Scott *et al.*, 2012; Khatchikian *et al.*, 2015). Some bird migrations can carry ticks to northern Ontario (including arctic ecoregions), beyond suitable living conditions for ticks (Ogden *et al.*, 2015). However, it is possible that as the climate warms, these areas will become adequate habitat and the ticks' range will further expand. The rate of range expansion for blacklegged ticks has been predicted to be approximately 46 km/year, with the rate of tick

colonization being slower and more variable (Leighton *et al.*, 2012; Clow *et al.*, 2017). The majority of the southern Ontario landscape is composed of deciduous ecosystems, and therefore suitable habitat for blacklegged ticks (Ogden *et al.*, 2006). As mean temperatures increase due to climate change, more of northern Ontario's terrain will become similar to the southern landscape and will be suitable for this species to inhabit (Ogden *et al.*, 2006; Leighton *et al.*, 2012).

Leighton *et al.* (2012) suggested that by 2030 blacklegged tick populations will be established across Ontario, exceeding beyond the current geographical range (Public Health Ontario 2019; Figure 0.3). This expansion will potentially lead to new interactions between blacklegged ticks and populations of deer mice that have never interacted with them before (*i.e.* naïve to this interaction), leading to the spread of multiple infections that can be transmitted by these ticks.

The wood tick (*Dermacentor variabilis*) is another medically important species abundantly found in brush field and forest habitats across the eastern United States and in several regions in Canada, including southern Ontario (Sonenshine, 2018; Minigan *et al.*, 2018). There are many ecological similarities between blacklegged ticks and wood ticks. *Peromyscus* mice are fundamental for the development of the larval and nymph stages of this species (Sonenshine, 2018), much like they are for blacklegged ticks (Ostfeld *et al.*, 1996; Estrada-Peña & de la Fuente, 2014). They have been found in similar habitats (Morshed *et al.*, 2003; J Curtis & E Fellin, pers. obs.), and are active at similar times of year (between May and September; blacklegged ticks; Ferreri *et al.*, 2014; wood ticks; Minigan *et al.*, 2018), although wood ticks have been found in areas farther north than blacklegged ticks (Figure 0.3: Public Health Ontario, 2019; Figure 0.4: Minigan *et al.*, 2018). The wood tick's range is also expanding with climate change and is expected to have a northwards expansion as more habitat becomes suitable (Minigan *et al.*, 2018). Wood ticks are the primary vector for the bacterium *Rickettsia rickettsii*,

which can cause Rocky Mountain spotted fever in humans and domestic animals, although it is also capable of transmitting *Coxiella burnetii* and *Francisella tularensis* at extremely low frequencies in Ontario (Wood *et al.*, 2016). It is expected, however, that as this tick's range expands, their population dynamics will change, altering the exposure of the pathogens they carry to humans and other animals (Minigan *et al.*, 2018).

## **Study Sites**

The sites chosen for this research overlap with the progression of the geographic expansion of blacklegged ticks and wood ticks into Ontario from both its southern and eastern populations (Hamer *et al.*, 2014; Clow *et al.*, 2016; 2017; Minigan *et al.*, 2018), and are based off of range expansion predictions of Leighton *et al.* (2012), Ogden *et al.* (2013) and Minigan *et al.* (2018; Figure 0.4). Three different sites in Ontario were visited for data and specimen collection. Each specific location was chosen to encompass the different levels of tick establishment. The most southern site (Long Point Provincial Park) has well established populations of blacklegged ticks and wood ticks (Watson & Anderson, 1976; Lindsay *et al.*, 1991). The other sites (Queen Elizabeth II Wildlands Provincial Park and Algonquin Provincial Park) are areas with no established populations of blacklegged ticks, although Queen Elizabeth II Wildlands Provincial Park is geographically close to Orillia, Ontario, which is a documented area where blacklegged ticks are prevalent (Public Health Ontario, 2019), and has known established American dog or wood tick (*Dermacentor variabilis*) populations (Minigan *et al.*, 2018; P. Careless, 2019, pers. comm.). All three areas are within the geographical habitat range of deer mice (Figure 0.1: Bedford and Hoekstra, 2015). I travelled from the most southern field

site (Long Point Provincial Park) northwards towards Algonquin Provincial Park, up a latitudinal gradient (Figure 0.5).

Long Point Provincial Park (LP; 42.5817° N, -80.3952° W) is a small park, covering a land area of just 1.50 km<sup>2</sup> and located in Port Rowan, Ontario (Long Point, 2019). LP hosts the oldest known established blacklegged tick population in Ontario (Watson & Anderson, 1976). The study site in Long Point was approximately 1.11 km<sup>2</sup> (Figure 0.5), encapsulating the forest space of the entire park. It should be noted that Long Point has been faced with seasonal flooding for several decades (Rasid *et al.*, 1992), and during this field season parts of the forest floor were submerged in water, providing a higher *Peromyscus* population density in the small land area than expected.

Queen Elizabeth II Wildlands Provincial Park (QEW; 44.7534° N, -78.7844° W) is located in Kawartha Lakes, near Orillia, Ontario. It is 335.05 km<sup>2</sup> (Queen Elizabeth II Wildlands, 2019). Leighton *et al.* (2012) predicted this area to have blacklegged tick populations establish by 2020. As of 2019, Orillia is the northernmost area in southwestern Ontario where there is a risk of Lyme's disease due to blacklegged tick presence (Public Health Ontario, 2019). Despite a large land area size, a small sample area in the southern region was used as a study site, similar to Long Point (Figure 0.5). Although blacklegged ticks are not prevalent here, this site offers a unique circumstance where there is a high prevalence of wood ticks (P. Careless, 2019, pers. comm.).

Algonquin Provincial Park (AP; 45.3402° N, -78.2618° W) is located between Georgian Bay and the Otter River in Ontario. Algonquin is Ontario's largest green space, bridging the gap between Southern and Northern Ontario, where ticks are expected to establish populations by 2022 (Leighton *et al.*, 2012). Currently, there are no documented sightings of either wood ticks

or blacklegged ticks in Algonquin, and so the Algonquin population of deer mice acted as a control group. The land area of AP is 7723 km<sup>2</sup> (Algonquin, 2019), however, I conducted my research in a smaller concentrated area at the Algonquin Wildlife Research Station (WRS). There are established traplines set up at the WRS for the long-term small mammal study that I used to collect data and specimens (Falls, 1953; Fryxell *et al.* 1998; Falls *et al.*, 2007; Bobbie *et al.*, 2016). During my data collection, these traplines were exclusively used for my study, despite typically being included in the long-term study. However, no mice found at this site were ear-tagged from previous studies, indicating these were all new individuals, avoiding any potential bias in capture effort or sampling as these mice had not been handled before. The ectoparasite communities on these mice were used as a comparison to determine any differences in parasite community assemblage when ticks are present or absent. The traplines here were set farther apart from each other than the other field sites (Figure 0.5), but were all 90 m long, so the overall trapping effort was the same.

Although it is not entirely clear when wood ticks initially established in LP or QEW, wood ticks have been found in Ontario “south of 44 degrees latitude” since the 1960s (Wilkinson, 1967), and the first documentation (according to Public Health Ontario) of wood ticks in LP was by Barker *et al.* (1988). Wood ticks have also been considered a common species in Canada since 1967, whereas blacklegged ticks were considered non-endemic at this time (Scholten, 1977), with their first establishment being documented in LP in 1976 (Watson & Anderson, 1976). There is no published data of the first record of wood ticks in QEW, although they are prevalent. Blacklegged ticks were also expected to be in this area (Leighton *et al.*, 2012), although none were found in this study. It can be concluded, however, that wood ticks established in LP before QEW, due to the northward expansion of both species (Clow *et al.*,

2017; Minigan *et al.*, 2018). While other tick species, such as the squirrel tick (*Ixodes angustus*), have been found in AP at low abundance and prevalence, neither blacklegged ticks nor wood ticks have established this far north.

It is also important to note the difference in habitat between the northern and southern sites. Southern sites are deciduous, and the landscape composition gradually changes to coniferous, moving northwards (Ontario Biodiversity Council, 2015). In the south, the forested areas where tick populations are established tend to be smaller provincial or national parks, between dense urban settlements. In northern Ontario, there tend to be larger landscapes dedicated as conservation territories, allowing for greater unfragmented areas where tick hosts can move around more easily. In general, the prevalence of blacklegged ticks has been shown to be greater in larger land fragments compared to small areas (Wilder & Meikle, 2004). Although all sites are a part of different ecoregions with somewhat differing habitats (Figure 0.6), LP and QEW are from the same ecozone (Mixedwood Plains). Similarly, despite AP being in the Canadian Shield ecozone, this site is in the south of this ecozone, and close in proximity to QEW. Although there are differences in habitats across these sites, the flora is similar, having mixed forests of coniferous and deciduous trees (Crins *et al.*, 2009). Trap sites were set in areas of mixed wood forests at each location, having both deciduous and coniferous trees, as well as shrubs, ferns, and both dense and sparse canopy cover in order to have as similar habitat as possible across sites that differed in ecozones.

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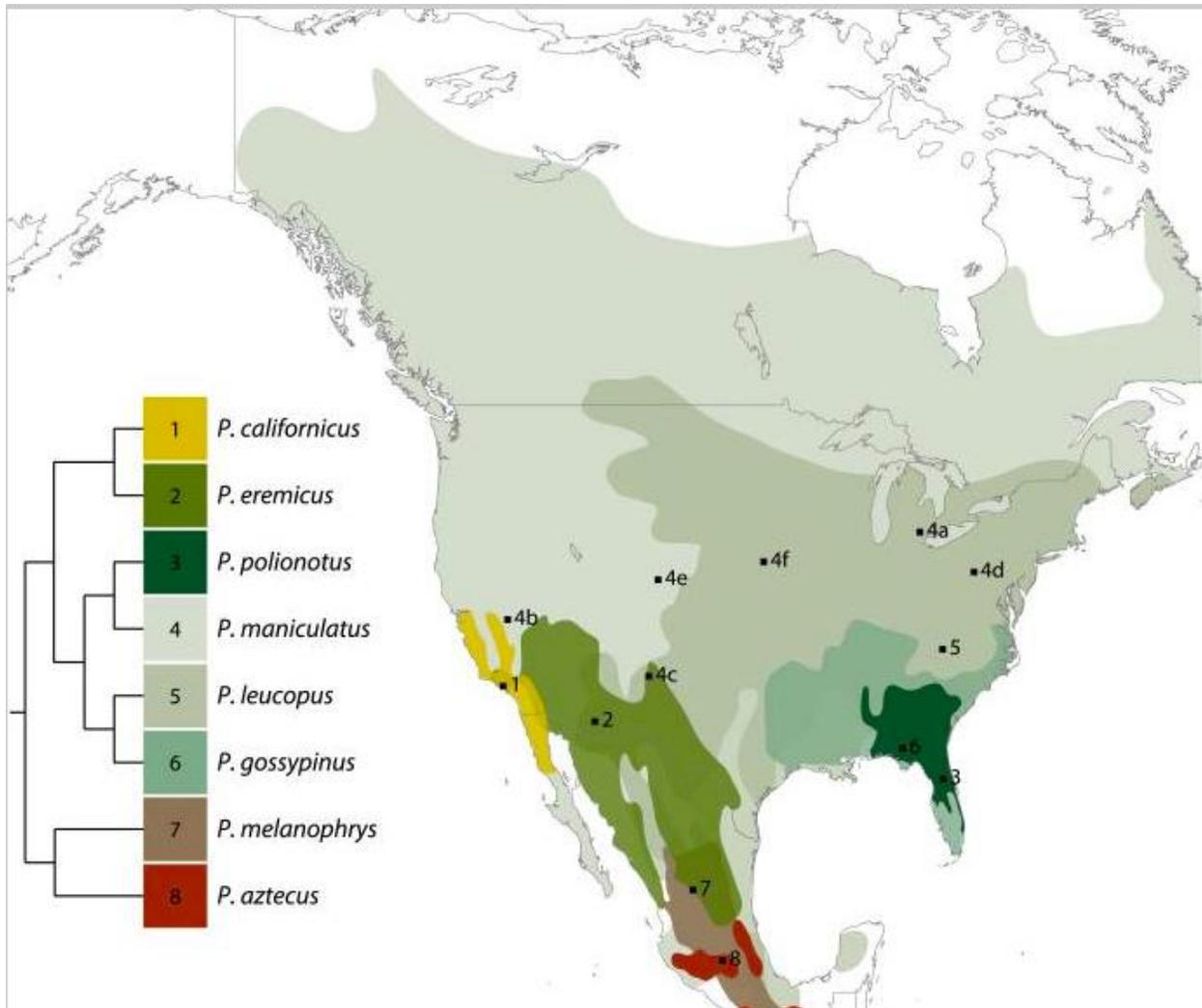
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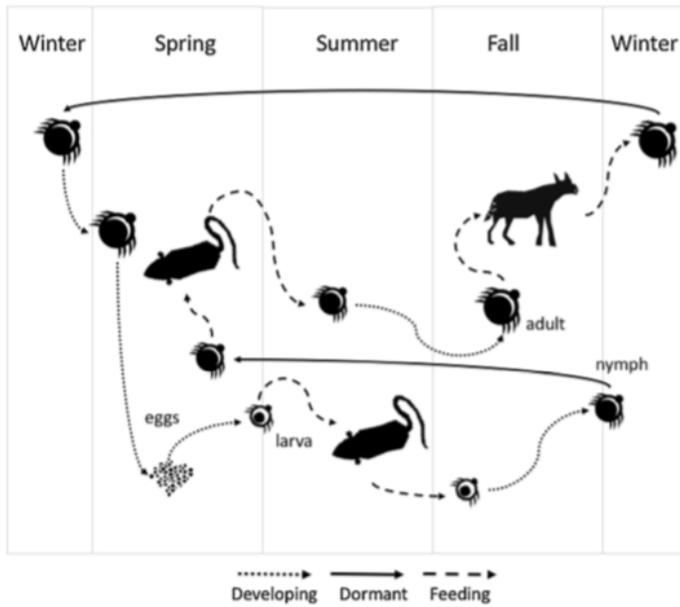
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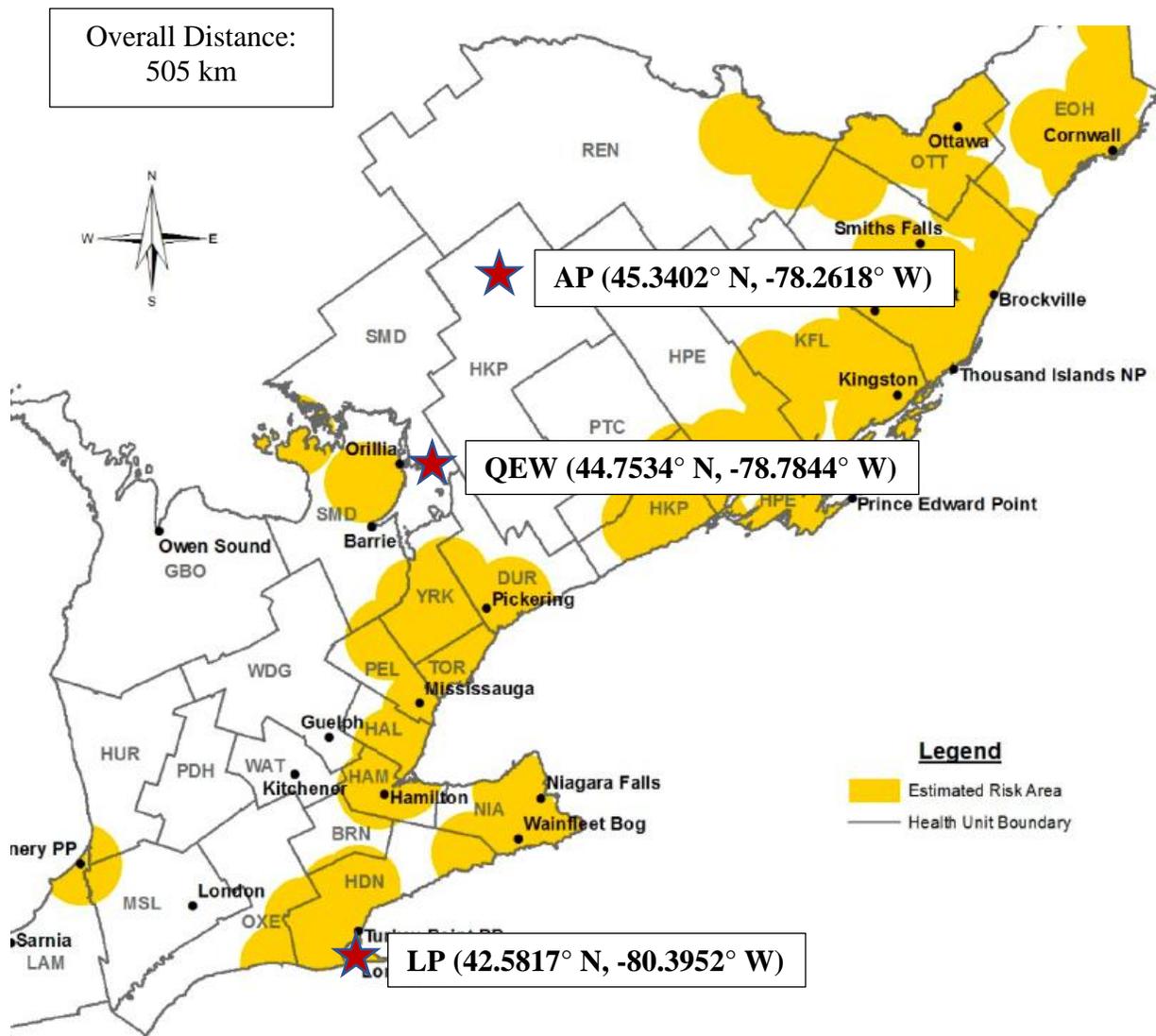
## General Figures and Tables



**Figure 0.1.** North American ranges of all *Peromyscus* species found in North America. The deer mouse (*Peromyscus maniculatus*) is the most abundant species of this genus, and as this figure shows, it also has the most extensive range (Bedford and Hoekstra, 2015).



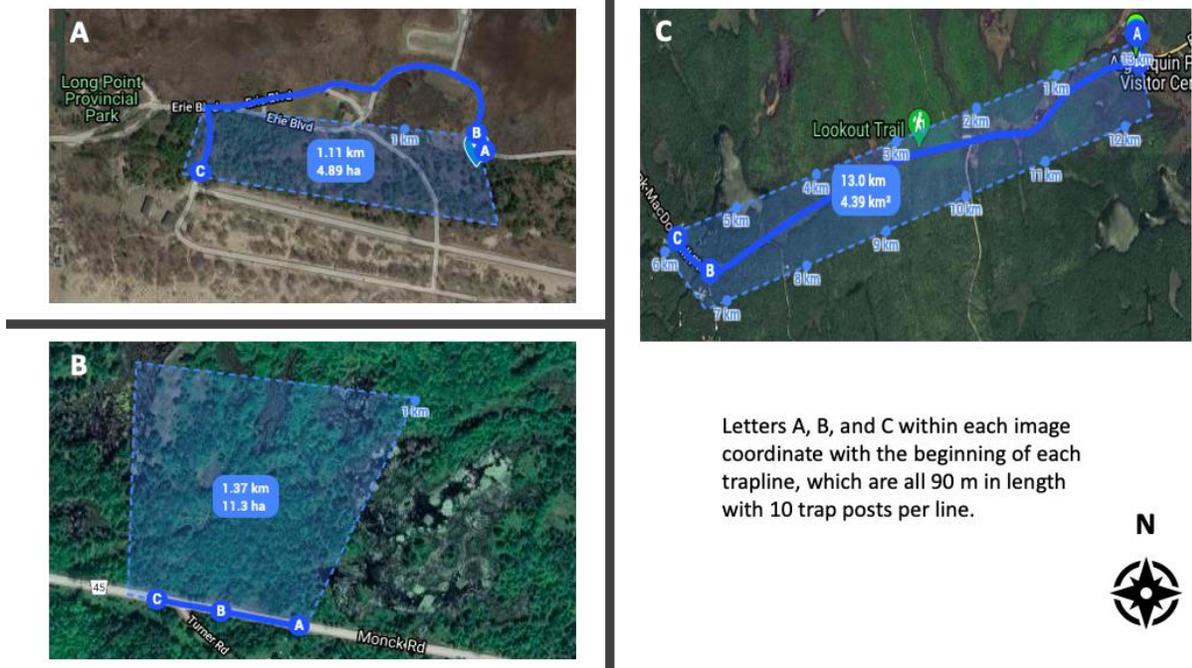
**Figure 0.2.** The life cycle of blacklegged ticks (*Ixodes scapularis*), illustrating the different developmental stages throughout seasons (Halsey & Miller, 2018).



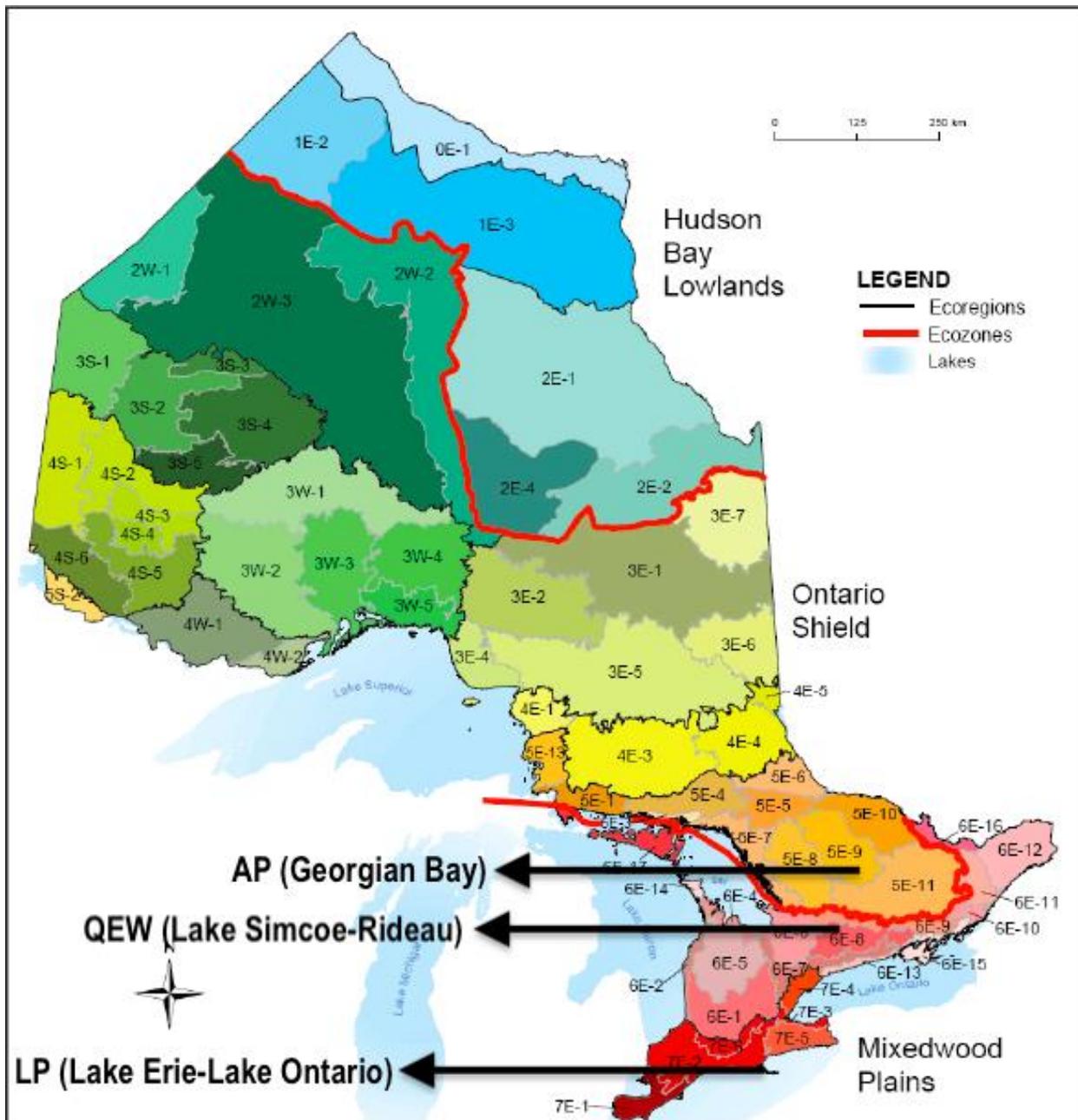
**Figure 0.3.** The estimated risk areas in Ontario, Canada where people are most likely to come in contact with blacklegged ticks, based on passive and active tick surveillance (Public Health Ontario, 2019). Red stars indicate sampling site locations.



**Figure 0.4.** The current distribution of suitable habitat for wood ticks in North America, highlighting the area of Ontario, Canada where the study is focused. The darker grey an area is, the better suited habitat that space is for wood ticks (Minigan *et al.*, 2018).



**Figure 0.5.** Land area of field sites where traplines were set and sampling occurred in Ontario, Canada. A) field Site at Long Point Provincial Park, B) field site at Queen Elizabeth II Wildlands Provincial Park, and C) field site at Algonquin Provincial Park (Google Maps, 2020).



**Figure 0.6.** Map of the ecoregions (delineated by black lines) and ecozones (delineated by red lines) of Ontario, Canada, including the ecoregions and ecozones for each site, depicting the similarities and differences across locations (Crins *et al.*, 2009). Black arrows indicate where sampling sites are and their respective ecoregions (in parentheses).

**Chapter I:**

**Effects of Ixodid Tick**

**Infestation on Hemoglobin**

**Levels of *Peromyscus***

***maniculatus***

## **Abstract:**

Deer mice (*Peromyscus maniculatus*) are hosts to several ixodid tick species as well as the associated tick-borne pathogens they can spread. As tick expansion continues northwards in Ontario, Canada, naïve host populations of deer mice are likely to become infested by ticks and susceptible to the physiological effects that ticks can have on them via blood-feeding. Prevalence of hematophagous ticks, such as blacklegged ticks (*Ixodes scapularis*) and wood ticks (*Dermacentor variabilis*), can affect hemoglobin levels of the deer mice they infest. Hemoglobin levels were compared and analyzed in deer mice populations at three different sites with varying exposures to ticks. It was expected that hemoglobin levels would significantly differ depending on how long ticks have been established at the sites and that a mouse population that was naïve to ticks would have the highest hemoglobin levels. My results suggested that the abundance of blacklegged and wood ticks on individual mice had the most significant negative effect on the hosts' hemoglobin levels, but the interaction between these two species did not. Average hemoglobin levels between populations also varied, where there was a significant difference between the source population with the longest established tick populations and the source population where neither blacklegged nor wood ticks were prevalent. As the ticks' ranges expand and they become more abundant, it is important to understand how their prevalence and intensity can alter the physiology of their hosts, potentially affecting their own range expansion and the spread of the diseases they may carry.

**Key Words:** *Dermacentor variabilis*, hemoglobin, *Ixodes scapularis*, parasite-host interactions, *Peromyscus maniculatus*

## Introduction

Hematophagous ectoparasites can negatively impact vertebrate hosts via blood-feeding (Bordes & Morand, 2011; Godinho *et al.*, 2013) and the physiological effects of these parasites on hosts have been widely studied (Dryden *et al.*, 1991; Carleton, 2008; Pfäffle *et al.*, 2009; Hersh *et al.*, 2014; Jones *et al.*, 2019). Blood-feeding by ectoparasites is known to be detrimental to the physiology of their hosts and can even prove fatal in some cases. This includes reduced hemoglobin levels (Carleton, 2008), regenerative anemia (Pfäffle *et al.*, 2009) and blood-loss related mortality (Jones *et al.*, 2019). Blood-feeding is important for ectoparasites, such as hard-bodied ticks (Acari: Ixodidae), because they require blood meals to reach subsequent life stages (Arsnoe *et al.*, 2015).

As the global climate warms and more habitat becomes suitable for ixodid tick species, the geographic ranges of these species expand into new areas (Leighton *et al.*, 2012; Clow *et al.*, 2016; 2017; Minigan *et al.*, 2018) where they can potentially establish and infest host populations previously naïve to such interactions. Variation in parasite susceptibility between naturally exposed and naïve populations has mostly focused on internal parasites (Brockhurst *et al.*, 2007; Hasu *et al.*, 2009; Sheath *et al.*, 2018). On an individual level, cattle calves (*Bos taurus*) infested with cattle ticks (*Rhipicephalus microplus*) have exhibited a significant reduction in hematological parameters, including hemoglobin when compared with non-infested calves (Kaur *et al.*, 2017). However, little is known about the effect of tick infestations on naïve mammal host populations, despite the geographic expansion of these pathogen vectors (Rand *et al.*, 1993; Larson *et al.*, 2018; Sonenshine, 2018). In general, naïve hosts are especially vulnerable to initial parasitic interactions (Brockhurst *et al.*, 2007). Contrastingly, hosts that have prior exposure to ectoparasites tend to be more resistant than hosts that have not been exposed to

them, due to their adaptive responses to promote stronger immune defences (Kennedy, 2010; Jones *et al.*, 2015).

Local adaption has been studied between parasite and host, but there is a general gap in literature on the relationship between hematophagous parasites and the effects of their blood-feeding on their hosts (Kutzer & Armitage, 2016; Papkou *et al.*, 2016). Several blacklegged ticks (*Ixodes scapularis*) can feed from white-footed mice at once or in succession due to exposed skin lesions that allow ticks to infest the same host despite a strong inflammatory response at the bite sites that can increase host resistance (Anderson *et al.*, 2017). Yet, insufficient immune responses during developmental periods could result in mice being vulnerable to other bacterial infections (Rosales *et al.*, 1999; Dlugosz *et al.*, 2014).

Blacklegged ticks and wood ticks (*Dermacentor variabilis*) tend to feed on *Peromyscus* mice at immature stages (Ostfeld *et al.*, 1996; Sonenshine, 2018), and have been documented sharing the same areas on a host (micro-habitat; Morshed *et al.*, 2003; Shaw *et al.*, 2003; Gómez-Rodríguez *et al.*, 2015). Although white-footed mice tend to be the preferred host of blacklegged ticks, and deer mice (*P. maniculatus*) have been found to have lower rates of tick infestations, both host species are equally competent vectors of tick-borne pathogens (*i.e.*, *Borrelia burgdorferi* and *Anaplasma phagocytophilum*: Rand *et al.*, 1993; Larson *et al.*, 2018). However, since ticks feed on mice for a short period of time, it is difficult to measure long-term impacts that they may have on host fitness. There is evidence that immature blacklegged ticks may not affect mouse survival in established host populations (Hersh *et al.*, 2014), although there are no comparative studies between experienced host populations that have cohabited with an established tick population versus inexperienced (naïve) ones.

Other ectoparasites that feed on *Peromyscus* mice for long periods of time (*i.e.* fleas, mites) are consistently affecting the vulnerability of their host's immune response (Mize *et al.*, 2011), reducing hemoglobin levels due to constant blood loss (O'Brien *et al.*, 2003), which could cause an insufficient production of hemoglobin, and therefore iron (Judy & Price, 1958; Andrews, 1997). Black rats (*Rattus rattus*) that were iron-deficient were found to be susceptible to infections of the pathogen *Salmonella typhimurium* (Baggs & Miller, 1973). It is possible that similar results may be found in other rodents. Investigating how ectoparasite feeding habits influence the hematology of deer mouse hosts – with and without the presence of ticks – will allow us to better understand how blacklegged and wood ticks may impact northern mouse populations as these ectoparasites move northward (Leighton *et al.*, 2012).

The objective of this study was to determine if ticks have a significant effect on the hematology of deer mice by examining how tick prevalence and intensity influence the host's hemoglobin levels. Such data may provide insight into how ticks can affect the overall fitness of their hosts, and how these hosts adapt to novel parasites. I hypothesized that: 1) if mice are parasitized by a greater intensity of ticks, then they will have lower hemoglobin levels compared to mice infested with ticks at low intensities because more blood-feeding is occurring at one time and 2) deer mice that live in areas where ticks are prevalent should have lower hemoglobin levels than deer mice living in unestablished areas because they are being affected by blood-feeding. Individual mice that are infested with higher abundances of ticks are expected to have lower hemoglobin levels compared to a mouse with low tick abundance as more ticks are feeding from the same source at one time. Further, mice that inhabit areas where ticks are prevalent and at high intensities are expected to have lower hemoglobin levels than mice that are ecologically naïve to these parasites. Examining the differences between naïve and experienced mouse

populations as well as between individual mice can bridge the gap in our understanding of tick effects on a host and help make better predictions on how hematology can affect the ecology of *Peromyscus* mice.

## **Methods**

All methods in this study were approved by the Animal Care Committee (ACC) at Laurentian University, file number 6017269.

### *Field Methods*

Three sites in Ontario, Canada were visited for data and specimen collection from May-August 2019. The most southern site, Long Point Provincial Park (LP; 42.5817° N, -80.3952° W) is an area where tick populations have been long-established (Watson & Anderson, 1976; Lindsay *et al.*, 1991). The other two sites, Queen Elizabeth II Wildlands Provincial Park (QEW; 44.7534° N, -78.7844° W) and Algonquin Provincial Park (AP; 45.3402° N, -78.2618° W) are areas with no established populations of blacklegged ticks, although QEW has known established wood tick populations (Minigan *et al.*, 2018; P. Careless, 2019, pers. comm.).

Neither blacklegged ticks nor wood ticks have colonized AP (Public Health Ontario, 2019). The sites chosen for this experiment convey the progression of the geographic expansion of blacklegged ticks and wood ticks in Ontario from both its southern and eastern populations (Hamer *et al.*, 2014; Clow *et al.*, 2016; 2017; Minigan *et al.*, 2018), and are areas within the geographical habitat range of deer mice (Bedford and Hoekstra, 2015). Traplines were placed in similar forest types across sites to maintain a consistent or similar habitat, although sites were from different ecozones (Crins *et al.*, 2009).

At each site, three traplines at least 0.3 km apart from each other were set up with twenty Longworth traps (Penlon Ltd., Oxford, U.K) set per line. Traplines were 90 m long and had two Longworth traps 10 m apart from each other every 10 m (Fryxell *et al.*, 1998; Falls *et al.*, 2007). Traps were baited with water-soaked sunflower seeds and were set half an hour before sunset (2000h-2100h). Deer mice are nocturnal, and so traps were checked half an hour before sunrise (0430h-0530h) after the mice had been active through the night (Clark & Durden, 2002). The traps were set five days each week. They were checked three days consecutively, with a one-day break, and then checked for two more consecutive days to reduce the amount of stress mice may have from repeated captivity or trap response behaviour (Nichols *et al.*, 1984). To account for change in seasonality, and the difference in parasite assemblages that occurs within the spring-summer months, I alternated between the three sites in ascending latitude, spending one week at each site, three times, for a total of three weeks per site. By alternating sites within the season, I was able to account for the different assemblages that may be occurring within populations, as temperatures change and specific parasites occur in different quantities, therefore removing potential seasonality bias to my analysis for parasite community assemblages.

Mice caught in traps were removed from traps and weighed using a Pesola® scale ( $\pm 0.1$ g). Age of mice was determined by body mass, where mice that were 15 g or less were considered juveniles, while mice greater than 15 g were considered adults (Banfield, 1974; Schmidt *et al.*, 2019). Sex and reproductive status were determined visually. Enlarged testes in males and perforated vagina and presence of nipples in females indicated reproductive individuals (Gaitan & Millien, 2016). No pregnant females were included in this study. Age of ticks was determined by the number of legs the tick had, where larva have six legs, and nymphs have eight (Lindquist *et al.*, 2016). Julian date was also recorded.

While mice were collected from the traps at each site, each mouse was examined for 60 seconds for sampling of ectoparasites (Patterson *et al.*, 2013). Individual arthropod specimens were combed off the host using a louse comb sterilized with ethanol (Hawlena *et al.*, 2006; Patterson *et al.*, 2013) or plucked off, using sterilized tweezers (Bobbie *et al.*, 2016). These specimens were then placed in a collection vial with 80% ethanol (Krogmann & Holstein, 2010). Tick identification was completed using guides (Lindquist *et al.*, 2016; Dubie *et al.*, 2017), and were later confirmed by Dr. Robbin Lindsay from the National Microbiology Laboratory (Winnipeg, Manitoba, Canada). Similarly, fleas were identified using an identification key (Holland, 1985), and later confirmed by Dr. Terry Galloway from the University of Manitoba. Two unique mite groups from the Trombiculidae family were identified by Dr. Heather Proctor (University of Alberta).

Deer mice were distinguished from white-footed mice by morphological characteristics. This included differences in ear length (mm; Kamler *et al.*, 1998), tail bi-colouration, tail “penciling” (J. Bowman, 2019, pers. comm.), and degree of dorsoventral colouration (Buchholz & Dick, 2017). Captured mice were placed in a 50 mL Falcon conical centrifuge tube with a hole in the tip to allow for the mouse to breathe. By pushing gently above the tibia, the right hind leg of the mouse could be extended out of the tube and shaved using an electric trimmer (Parasuraman *et al.*, 2010). Needles (20G) were used with a 1 mL syringe in accordance with Standard Operating Procedure for Laurentian University and general principles of laboratory blood collection (Parasuraman *et al.*, 2010). Blood samples consisted of 0.007 mL of blood/g of the mouse’s body mass taken from the saphenous vein in the shaved hind leg of the mouse via needled syringe (Randolph, 1980; National Research Council Institute for Laboratory Animal Research, 1996; PREDICT One Health Consortium, 2016). On site, the blood samples were

placed in a microcuvette specialized for the handheld hemoglobin analyzer to test for hemoglobin concentrations: HemoCue Hb 201+ analyzer (HemoCue AB, Ängelholm, Sweden; Tufts *et al.*, 2013; Weldon *et al.*, 2015). Results were given immediately by the analyzer and recorded.

It was necessary to shave the legs of the mice in order to acquire blood samples and so the shaved areas were also used as markers on the mice to indicate that they were already sampled to prevent sampling an individual more than once (Powell & Proulx, 2003). Although it is common for small mammals to be given ear tags for identification purposes in field studies (Harper & Austad, 2001; Hersh *et al.*, 2014; Torre *et al.*, 2016; Buchholz & Dick, 2017), Ostfeld *et al.* (1993) found that this method will increase tick infestations on the host's ears. It should be noted that the exposed shaved legs of the mice in this study were never observed to have ticks or bite marks in this specific area, implying shaving the legs of the mice did not result in the same bias as ear tags do in attracting ticks. Instead, because their right legs were shaved, it was clear which mice were recaptures when returning to each site every three weeks, as the fur on these mice had not fully regrown. These mice were documented as recaptures and were not repeatedly included in any statistical analysis as multiple blood samplings taken from one individual could provide lower hemoglobin levels over time (Weixelbaumer *et al.*, 2010), biasing the study. The legs of these recaptures were re-shaved, however, so that upon returning to the site in the next cycle, they could be clearly identified again.

### *Statistical Methods*

Statistical analyses were conducted in the R environment (R core Team 2019 [v3.6.1]) using base R and the package 'ggplot2' (Wickham, 2016 [v3.2.1]) for figures and linear models.

Prevalence and mean intensity of ectoparasites were completed using QPweb (Reiczigel *et al.*, 2019), where prevalence refers to the percentage of mice in a population with one or more ectoparasites (Bush *et al.*, 1997). For the purpose of this study, mean intensity refers to the mean number of ectoparasites found across mice in a population, whereas abundance refers to the number of ectoparasites found on individuals (Bush *et al.*, 1997). Shapiro-Wilks tests showed that hemoglobin levels were distributed normally across individuals ( $p = 0.09$ ).

To determine differences in hemoglobin levels among sites, a one-way ANOVA and Tukey's Honest Significance test were conducted, where hemoglobin levels were the response variable, and source population site (LP, QEW, AP) was the predictor. To account for possible variation across individuals in the ANOVA, a general linear model was performed first to see if there were any significant differences across host sex, host age, host body mass, and host population location (field site). In this model, only field site had a significant effect on hemoglobin levels (Table 1.1). Similar results were found in a general linear model with tick abundance (both blacklegged and wood ticks together) as the response variable (Table 1.2), suggesting that there is no difference in tick loads across host age, sex, or body mass; only field site. The effect of differences in tick prevalence on mouse population hemoglobin levels was measured by comparing mean hemoglobin levels between sites and across all individuals (LP+QEW+AP) when 1) all mice are included, 2) only mice infested with ticks are included, or 3) only mice not infested with ticks are included. Standard deviations of these means were also included. One-way ANOVAs were conducted for all individuals, and for each source population site comparing hemoglobin levels and tick prevalence (both blacklegged and wood tick together).

Julian date, population density per site (using Lincoln-Peterson index to determine population size via mark-recapture data; Grimm *et al.*, 2014), and engorgement level of ticks (determined visually, and categorized as 0 = not engorged, 1 = moderately engorged, 2 = substantially engorged; Hofhuis *et al.*, 2017) were variables initially considered. However, after performing Pearson correlation coefficient ( $r$ ) tests, they were found to be heavily correlated with other variables – Julian date and session ( $r = 0.92$ ), population density and location ( $r = -0.99$ ), tick engorgement level and wood tick abundance ( $r = 0.90$ ). The package ‘tidyverse’ (Wickham *et al.*, 2019 [v1.3.0]) was used to test variance inflation factors (VIF) for independent variables. To avoid severe multicollinearity, the variables with the highest VIF values were removed from analyses (Julian date, tick engorgement, and population density; Harrison *et al.*, 2018; Frost, 2019).

Across all individuals from all three sites, multiple general linear regressions were considered with hemoglobin levels as the response variable, comparing the effects of the predictor variables: number of wood ticks, number of blacklegged ticks, wood nymph presence (Y or N), presence of *Neotrombicula sp.* (Y or N) and presence of Trombiculidae *sp.* (Y or N) – separately and together (*Neotrombicula sp.* presence only, Trombiculidae *sp.* presence only, both *Neotrombicula sp.* and Trombiculidae *sp.* together), number of *O. leucopus*, sessions when each site was visited, age of mouse (adult or juvenile), sex of mouse (male or female), reproductive status of mouse (reproductive or non-reproductive), body mass of mouse, and source population (LP, QEW, AP). Sessions were divided by early (May-June), mid (June-July), and late (July-August) summer. Only two blacklegged nymphs were found, so only the presence of wood nymphs was measured.

Since the sample size for this study was small ( $n = 44$ ), Akaike information criterion correction (AICc) tests were conducted via R package ‘AICcmodavg’ (Mazerolle, 2019 [v2.2-2]). AICc analyses were used to determine linear models that provided the strongest explanation for affecting hemoglobin levels when  $\Delta\text{AICc}$  is  $< 3$ . Here all models within this threshold or ‘top model set’ were considered equally plausible (Harrison *et al.*, 2018). Pearson’s correlation formula was used to determine correlation coefficients between all predictor variables included in the top models and hemoglobin levels to determine effect sizes (Weber *et al.*, 2019). VIF multicollinearity tests were conducted on all multiple linear regression models where all independent variable VIF values were  $< 5$  (Harrison *et al.*, 2018; Frost, 2019). Source population was removed as a predictor, since it had  $\text{VIF} > 5$  (5.13) and was moderately correlated with blacklegged tick and wood tick abundances ( $r = -0.64$  and  $-0.54$ , respectively). This was expected, as the populations are from field sites that were chosen based on their differences in tick prevalence and intensity. Wood nymphs had the highest VIF (3.65), because it was correlated with wood tick abundance ( $r = 0.65$ ). All other variables had  $\text{VIF} < 3$ . Once the top model set was determined, multiple linear regression models were performed using the variables from these models.

## Results

A sample of 44 individual deer mice was included in analyses among all sites. This included 16 adult female mice and 3 juvenile female mice, 20 adult male mice, and 5 juvenile male mice. Dr. Terry Galloway identified one flea species, which was found across sites: *Orchopeas leucopus*. Dr. Heather Proctor was able to identify two separate groups from the mite specimens collected. One group was identified to the genus *Neotrombicula*. The other group was

determined to be taxonomically distinct enough to be a different species from the *Neotrombicula* specimens but could not be identified beyond family; Trombiculidae. For the purpose of this study, the mite groups will be defined as '*Neotrombicula sp.*' and '*Trombiculidae sp.*'. There were 111 wood ticks (16 of which were nymphs), 28 blacklegged ticks (2 of which were nymphs), and 15 *O. leucopus*. There were 15 instances of *Trombiculidae sp.* and 8 of *Neotrombicula sp.* on deer mice. The average population sizes for each site across sessions (early, mid, and late summer) were determined by the Lincoln-Peterson index: LP = 28 mice, QEW = 18 mice, AP = 11 mice. These results, along with the land area used for trap sites determined population density (Table 1.3). LP had the greatest intensity and prevalence of both focal tick species, while AP had the highest prevalence of *O. leucopus* and *Neotrombicula sp.*, and QEW had the highest prevalence of *Trombiculidae sp.* (Table 1.4).

Models with the lowest AICc scores determined which predictor variables may be affecting the response variable – deer mouse hemoglobin levels – most significantly. Across the top two models, the explanatory variables were the abundance of blacklegged ticks on an individual, the abundance of wood ticks on an individual, the interaction between these two species, and the prevalence of wood tick nymphs on an individual (Table 1.5). Multiple linear regressions were performed for the two models. Across the models, it was found that the abundance of blacklegged ticks and wood ticks significantly affected hemoglobin levels ( $p < 0.05$ ), but their interaction did not ( $p = 0.65$ ; Table 1.6; Figure 1.1). Pearson correlation coefficients between hemoglobin levels and tick species (both tick species together, blacklegged ticks, and wood ticks separately) were moderately and negatively correlated ( $r = -0.49, -0.51, -0.5$ , respectively; Table 1.7), suggesting moderate effect size (Weber *et al.*, 2019). Across the two models, the prevalence of wood nymphs was also found to significantly affect hemoglobin

levels, ( $p = 0.01$  and  $p = 0.03$ ; Table 1.6; Figure 1.2), however, the Pearson correlation value for wood nymphs was both negatively and weakly correlated to hemoglobin ( $-0.21$ ; Table 1.7), suggesting small effect size (Weber *et al.*, 2019). However, the estimates for wood nymph presence were positive (1.95, 1.85; Table 1.6), despite a negative relationship with hemoglobin levels (Figure 1.2). This could be due to collinearity between wood nymph presence and wood tick abundance. None of the top models included *Neotrombicula sp.* presence, Trombiculidae *sp.* presence, *O. leucopus* abundance, field sessions, host age, host sex, host reproductive status, host body mass, or source population (LP, QEW, AP), suggesting that these variables had no significant effect on hemoglobin levels.

An ANOVA testing the differences in hemoglobin levels across sites showed that there was a significant difference between hemoglobin levels depending on the location where the deer mice were found ( $p = 0.01$ ). A Tukey's Honest Significance test showed that AP hemoglobin levels were significantly higher than those at LP ( $p_{\text{adj}} = 0.01$ ), and the differences between LP and QEW hemoglobin levels and between QEW and AP were not significant ( $p_{\text{adj}} = 0.09$  and  $p_{\text{adj}} = 0.99$ , respectively; Figure 1.3). These differences are further explained by the differences in hemoglobin levels due to tick prevalence. ANOVA results showed a significant difference when ticks were present versus absent on a host across sites ( $p = 0.012$ ), although within sites, this difference was not significant: QEW ( $p = 0.492$ ), and AP ( $p = 0.814$ ; Table 1.8). Since LP mice were all infested, tick prevalence could not be compared, however, LP mice did have the lowest mean hemoglobin levels (Table 1.8).

## Discussion

### *Tick Abundance*

I found that blacklegged tick and wood tick abundances negatively affected hemoglobin levels in individual hosts. Most individual mice had hemoglobin levels much greater than 13.2 g/dL, which is the known average of lab-reared deer mice not infested with any ectoparasites (Wiedmeyer *et al.*, 2014), suggesting this number is not necessarily accurate for wild-caught mice, so it is unknown if the reduction in hemoglobin levels for the deer mice in this study is adversely affecting their physiology. However, the prevalence and intensity of ticks has been shown to have a negative effect on hemoglobin levels in other mammals. Two studies found that cattle calves infested with ticks had substantially lower hemoglobin levels compared to their non-infested counterparts (Rahman *et al.*, 2009; Kaur *et al.*, 2017). Similarly, a study on moose calves (*Alces alces*) that were moderately to severely infested with winter ticks (*Dermacentor albipictus*) had high mortality rates (Jones *et al.*, 2019). Results from my study also show that tick prevalence and intensity affect hemoglobin levels in deer mice, which could be detrimental to their physiology. In European hedgehogs (*Erinaceus europaeus*), the blood-feeding activity of two tick species of the genus *Ixodes* (*I. ricinus* and *I. hexagonus*) resulted in lower hemoglobin levels and led to regenerative anemia in several individuals (Pfäffle *et al.*, 2009). Gaitan & Millien (2016) found that white-footed mice infested with higher intensities of blacklegged ticks tended to have lower movement rates than mice hosting fewer ticks, suggesting that higher infestation rates incapacitate these mice in some way. This may be related to lower hemoglobin levels, which can reduce motor skills (Nicolas *et al.*, 2002). These studies suggest that the negative impacts of blacklegged ticks and wood ticks could be detrimental, although not necessarily fatal. Hersh *et al.* (2014) found that larval tick burdens do not affect the survival of

*Peromyscus* hosts, and the majority of the ticks found in LP and QEW were found at the larval stage. Since larval ticks cannot transmit pathogens, but rather acquire them during this life stage (and then pass them on as nymphs; Huang *et al.*, 2019), Hersh *et al.*'s (2014) study suggests that it is not the tick burden alone which is affecting their hosts.

Since blacklegged ticks were only found in LP, the mice in LP are the only ones whose hemoglobin levels are affected by these ticks. However, the prevalence and intensity of wood ticks at this site was much higher and so it could be that tick abundances and mouse population sources are confounding variables. Although one of the top AICc models suggested that the interaction between wood ticks and blacklegged ticks should also be a variable included in this relationship (Table 1.5), the interaction was not significant ( $p = 0.65$ ; Table 1.6). This suggests that each species is affecting their hosts independently rather than jointly, where there are no confounding effects. Moreover, despite both species having a significant negative effect on hemoglobin levels mice, the abundances of blacklegged ticks and wood ticks varied greatly. Infections can play a role in the hematology of the host, reducing blood cell counts, which can result in anemia (Westblade *et al.*, 2017). Some infections can affect a host's hematology, such as *Babesia* protozoans that invade mature red blood cells and directly affect hemoglobin concentrations (Borggraefe *et al.*, 2006). Blacklegged ticks can transmit *Babesia* protozoans to their *Peromyscus* hosts, but wood ticks cannot (Westblade *et al.*, 2017), so it may be possible that blacklegged ticks are indirectly affecting hemoglobin levels in hosts rather than directly by transmitting pathogens that affect the blood. A recent study found that *Babesia odocoilei* was identified in blacklegged ticks at Long Point (Milnes *et al.*, 2019), so it is possible this may be occurring in the focal host species, but this variable was not examined in this analysis.

My results did not find any differences in hemoglobin levels across host's sex. This is supported by several studies, where there was no difference found in hemoglobin levels between male and female deer mice (Wiedmeyer *et al.*, 2014; Eleftheriou *et al.*, 2020). Similarly, deer mouse hemoglobin levels did not differ across age or body mass. To my knowledge, no studies on hemoglobin levels differing with age or body mass have been conducted in mice or small mammals, but in humans age does not appear to influence hemoglobin levels (Raisinghani *et al.*, 2019). Similarly, hemoglobin levels in humans have not been found to be affected by differences in body mass index (Ghadiri-Anari *et al.*, 2014), however, tick burdens in *Peromyscus* mice are known to be associated with both body mass and sex (Dallas *et al.*, 2012).

### *Nymph Prevalence*

Wood tick nymphs were shown to have a significant effect in the given models. Since nymph ticks are larger than larval ticks and can feed from hosts for longer periods of time (Estrada-Peña & de la Fuente, 2014; Kocan *et al.*, 2015), they should consume more blood than their larval counterparts. Nymphs are also more likely to spread diseases, further affecting their host's physiology (Lindquist *et al.*, 2016). The larger blood meals and potential effects nymphs have on their host's physiology could explain how the prevalence of wood tick nymphs negatively affects hemoglobin levels. Yet, the majority of nymphs in this study were of this species only so the effects of blacklegged nymphs are unknown. Although there are few studies comparing the differences in blood meal volume between ticks in general, Koch and Sauer (1984) found that adult female wood ticks ingested greater volumes compared to blacklegged ticks. Although there has not been any comparison between larval and nymph ticks of these species, wood ticks tend to be larger than blacklegged ticks when unfed and engorged (Lindquist

*et al.*, 2016). It is thus possible that nymph wood ticks can acquire greater blood volumes than nymph blacklegged ticks. However, there was an imbalanced sample size between wood and blacklegged nymphs, so it could be simply that nymphs in general affect hemoglobin levels more than larval ticks in the study system.

### *Hemoglobin Variation Across Sites and Tick Prevalence*

It was expected that the variation in tick species prevalence between deer mouse populations would affect hemoglobin levels, such that deer mice from a population where ticks have not established would have higher hemoglobin levels than mice from a population where ticks have established. Within sites, it was expected that there would be a difference in hemoglobin levels depending on the tick species intensities. I found that hemoglobin levels significantly differed across sites, but not within sites. In particular, hemoglobin levels differed significantly between LP, which had the highest tick intensity and greatest tick prevalence, and AP, which did not have an established wood tick or blacklegged tick population. Within sites, there was not a significant difference in hemoglobin levels (Table 1.3). However, the sample sizes across sites were imbalanced, and QEW in particular had a small sample size ( $n = 8$ ), with only two infested hosts, while the one individual infested with a tick in AP, was not infested with a focal species. Due to the constraint in individual site sample size, results within each site may be misleading, and tick infestation effects could be hidden. Even so, my results suggest that in general, tick abundance can reduce a host's hemoglobin level and that tick intensity affects hematology. Within a population, however, hemoglobin levels are not significantly different between infested mice and non-infested mice. It is possible that by having prior exposure (or lacking exposure) to these ectoparasites in a population, the immune defences within the

population are similar (Nédélec *et al.*, 2016). This could result in non-significant differences in hemoglobin levels, regardless of whether or not ticks are prevalent on individual hosts in that population.

### *Limitations & Future Directions*

One variable not accounted for that could have an effect on hemoglobin levels in deer mice was the water intake of individual mice. Water-deprived laboratory mice that were restricted water for over 24 h, were documented to have significant variation in hematocrit, depending on the extent of their deprivation (Bekkevold *et al.*, 2013). Mice caught in traps in my study were not deprived of water, but it is possible that while trapped (or pre-trapping) individual mice may have been dehydrated for several hours. Other variables that were not included in this study include the microbiota of the host, which can affect hematology and can differ significantly across sites (Weldon *et al.*, 2015). Additional differences in sites must also be considered. Smaller land areas will have smaller mouse (and therefore smaller tick) populations. Therefore, negative impacts caused by tick infestations may hold a greater significance to the overall population in smaller populations, which have less genetic diversity and higher rates of disease transmission (King & Lively, 2012). This could be another reason why hemoglobin levels in LP were lower than the other two sites. Since LP is a much smaller site, having a land area of only 1.50 km<sup>2</sup>, the mouse population is not only smaller but denser than QEW or AP. The population density for LP is also inflated in this study, due to flooding in the area, which caused the mice to be more concentrated in dry areas where traps were placed (Wijnhoven *et al.*, 2005). Such environmental factors were not applicable in QEW or AP, which are already larger habitats. Sample size imbalance could also play a role, as only seven deer mice from QEW were

included in analyses. These mice also had low intensities of wood ticks, although some mice caught during this study in this area were found with high wood tick infestations, but hemoglobin samples were not taken so these individuals were not included in any analysis. Future research should focus on other factors that may play a role in parasite-host relationships, such as other co-infections and the microbiota, which can affect host physiology and hematology. To gain a better understanding of how naïve host populations may respond to the introduction of tick species to their ecosystem, experiments where these hosts are infected with ticks should be considered in a lab setting.

### *Conclusion*

Overall, hemoglobin levels differed at different degrees of tick abundance across individuals and prevalence across populations, although the most significant difference was between a population with a high infestation rate and a population with little to no infestation, suggesting that low infestation rates do not impact deer mice detrimentally on a hematological level. Tick abundance appeared to contribute most to hemoglobin differences on an individual level. When examining the effects of both tick species together, blacklegged ticks only, and wood ticks only, hemoglobin levels declined significantly when there were greater intensities of ticks on an individual. Despite blacklegged ticks being found at lower intensities than wood ticks, both species significantly affected hemoglobin levels. This may be due to the bacteria blacklegged ticks can carry or a difference in the host's resistance to blacklegged ticks relative to wood ticks, though further studies must be conducted to determine this. These results may suggest that deer mouse populations that will soon be introduced to blacklegged ticks may see their hemoglobin levels decrease more substantially compared to when wood ticks are introduced

to a new mouse population. The susceptibility to tick infestations and their effects has the potential to alter ecosystem processes that can ultimately affect other host species as zoonotic pathogens are transmitted from mice to ticks to other host species at higher trophic levels, including humans. Future studies should consider variables that were not measured in this project, including internal parasites and bacteria that the host may be infected with, as well as their gut microbiota, which can influence the overall biology of the host, including hemoglobin concentrations. More research on the interactions between co-occurring species should be done, considering the impacts of invasive introductions to naïve populations, as well as the co-evolution and local adaptations of parasites and hosts. This, along with experimental studies exposing naïve hosts to new parasites will help increase our understanding of host-parasite interactions.

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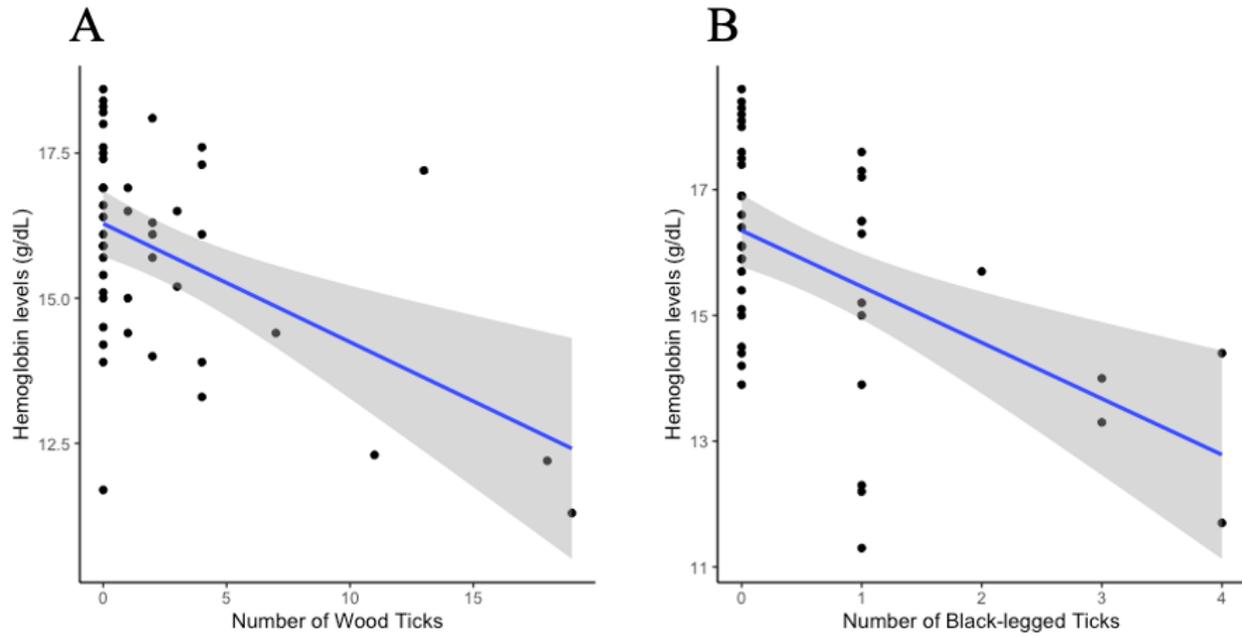
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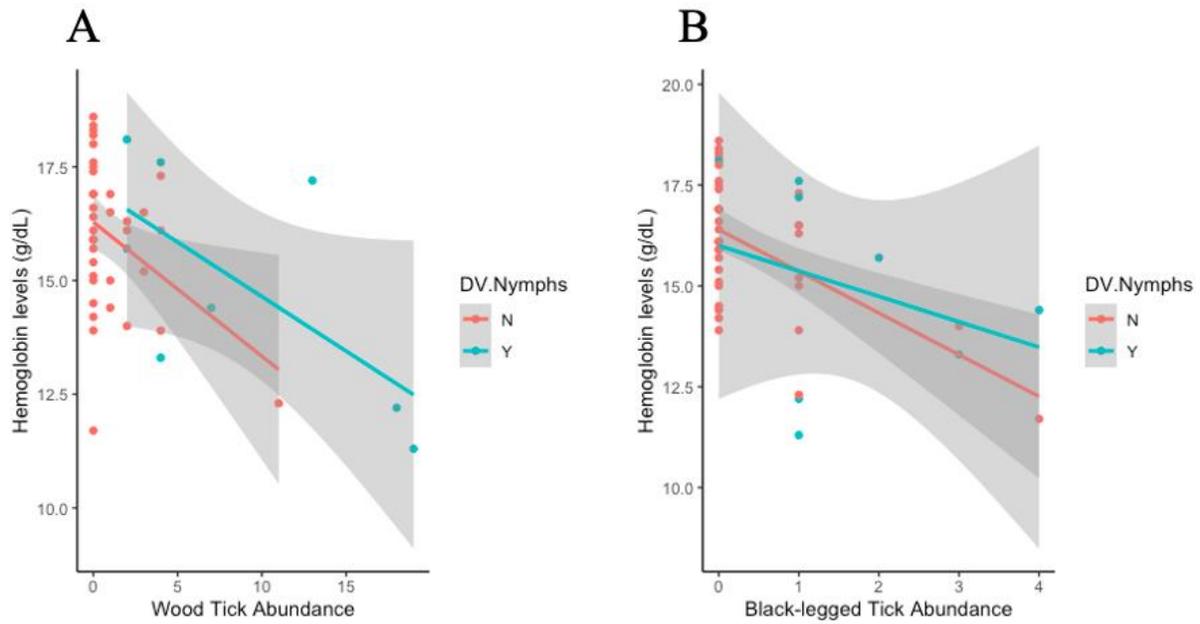
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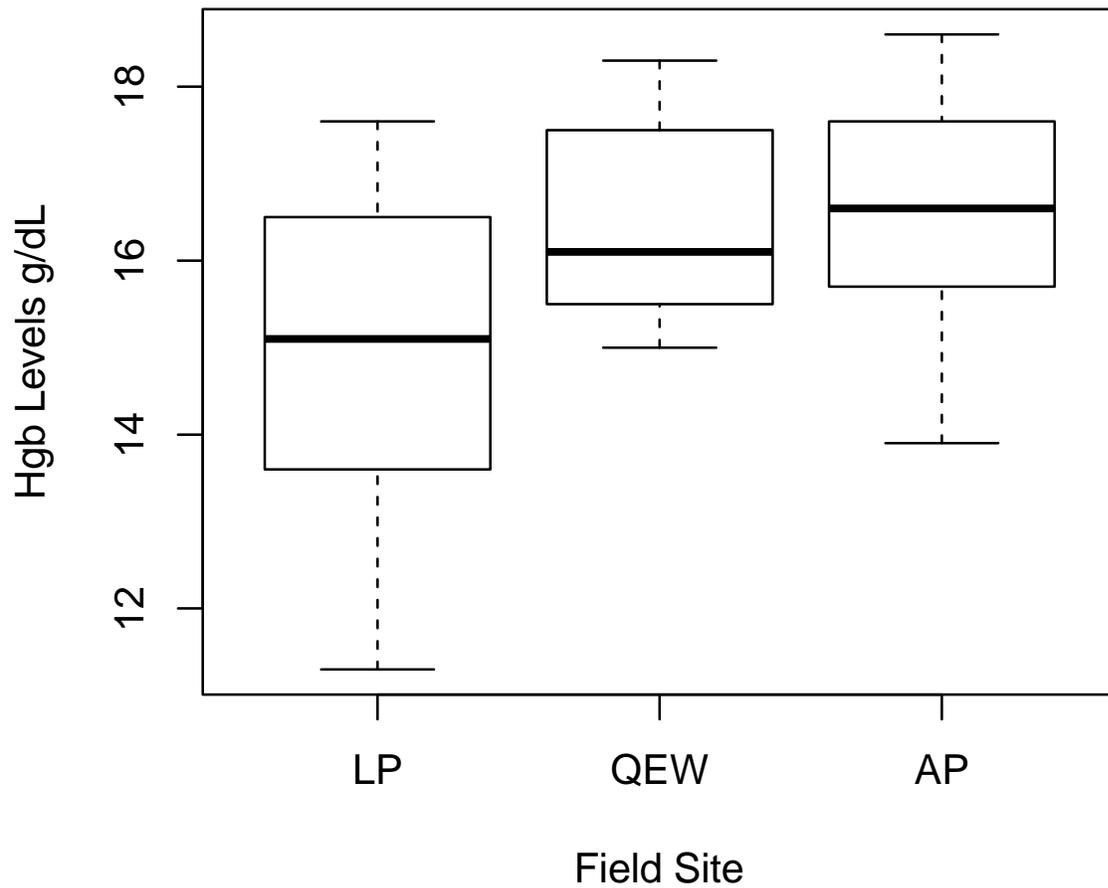
## Figures and Tables



**Figure 1.1.** General linear regression results depicting the effect of tick abundance on hemoglobin levels in deer mice (*Peromyscus maniculatus*;  $n = 44$ ). Each point indicates an individual host. Shaded area indicates the 95% confidence interval. A) Wood tick abundance is significantly affecting hemoglobin levels ( $p$ -value  $< 0.01$ ) as is B) blacklegged tick abundance ( $p$ -value  $< 0.01$ ; Table 1.5).



**Figure 1.2.** General linear regression results depicting the effect of tick abundance on hemoglobin levels in deer mice (*Peromyscus maniculatus*;  $n = 44$ ) when wood (*D. variabilis*) nymphs are present on the host. Blue points indicate that wood nymphs were prevalent on that individual host. Shaded area indicates the 95% confidence interval. A) Wood tick abundance was significantly affecting hemoglobin levels ( $p\text{-value} < 0.01$ ) as was B) blacklegged tick (*I. scapularis*) abundance ( $p\text{-value} < 0.01$ ), but their interaction was not ( $p\text{-value} = 0.65$ ). The presence of wood nymphs (DV Nymphs) was significant ( $p < 0.05$ ; Table 1.5).



**Figure 1.3.** Hemoglobin levels across each site (LP: n = 20. QEW: n = 7, AP: n = 17). ANOVA results show a significant difference across sites ( $p = 0.01$ ). Based on a Tukey's Honest Significance test the comparative differences between sites are LP-QEW ( $p \text{ adj} = 0.09$ ), LP-AP ( $p \text{ adj} = 0.01$ ), and QEW-AP ( $p \text{ adj} = 0.99$ ). LP = Long Point Provincial Park, QEW = Queen Elizabeth II Wildlands Provincial Park, AP = Algonquin Provincial Park.

**Table 1.1.** Results for general linear model examining the effects of deer mice (*Peromyscus maniculatus*) sex, age, body mass, and population (field site) on hemoglobin levels. This model had a residual standard error of 1.748 on 39 degrees of freedom. Multiple R-squared: 0.1973, Adjusted R-squared: 0.1149. Bolded values indicate variables where  $p < 0.05$ .

Variable	Estimate	Std. Error	t - value	Pr(> t )
Intercept	14.86	1.30	11.44	4.93e <sup>-14</sup>
Host sex	-0.35	0.54	-0.65	0.52
Host age	0.06	0.84	0.07	0.94
Host body mass	-0.01	0.06	-0.07	0.94
Field site	0.59	0.20	2.92	<b>0.006</b>

**Table 1.2.** Results for general linear model examining the effects of deer mice (*Peromyscus maniculatus*) sex, age, body mass, and population (field site) on general tick abundance (both blacklegged and wood ticks together). This model had a residual standard error of 4.331 on 39 degrees of freedom. Multiple R-squared: 0.6151, Adjusted R-squared: 0.5756. Bolded values indicate variables where  $p < 0.05$ .

Variable	Estimate	Std. Error	t - value	Pr(> t )
Intercept	11.32	3.22	3.52	0.001
Host sex	0.92	1.34	0.68	0.50
Host age	3.46	2.07	1.67	0.10
Host body mass	-0.06	0.14	-0.40	0.69
Field site	-3.80	0.51	-7.60	<b>3.28e<sup>-9</sup></b>

**Table 1.3.** Study sites, geographic coordinates, land area of parks and field sites, as well as average population size across (measured across sessions, determined by Lincoln-Peterson index) to determine population densities.

Sites	Latitude and Longitude	Total Land Area of Provincial Park	Land Area of Field Site	Average Population Size	Population Density per Field Site
Long Point Provincial Park (LP)	42.5817° N, -80.3952° W	1.50 km <sup>2</sup>	0.31 km <sup>2</sup>	28 mice	90 mice/km <sup>2</sup>
Queen Elizabeth II Wildlands Provincial Park (QEW)	44.7534° N, -78.7844° W	335.05 km <sup>2</sup>	0.47 km <sup>2</sup>	18 mice	38 mice/km <sup>2</sup>
Algonquin Provincial Park (AP)	45.3402° N, -78.2618° W	7723 km <sup>2</sup>	4.39 km <sup>2</sup>	11 mice	2.5 mice/km <sup>2</sup>

**Table 1.4.** Prevalence and mean intensity (average abundance) of ectoparasites found on deer mice (*Peromyscus maniculatus*) across all sites and per site. Numbers in parentheses indicate the sample size of deer mouse hosts for each row. Since mite groups were only measured as present (1) or absent (0) for all analysis, their intensities are unknown. LP = Long Point Provincial Park, QEW = Queen Elizabeth II Wildlands Provincial Park, AP = Algonquin Provincial Park.

	<i>I. scapularis</i>	<i>D. variabilis</i>	<i>O. leucopus</i>	Trombiculidae <i>sp.</i>	<i>Neotrombicula</i> <i>sp.</i>
All Sites (44)					
Prevalence	37.8%	48.9%	20.0%	33.3%	17.8%
Mean intensity	0.63 +/- 1.06	2.45 +/- 4.53	0.34 +/- 0.89	N/A	N/A
LP (20)					
Prevalence	85.0%	95.0%	10.0%	0.0%	5.0%
Mean intensity	1.40 +/- 1.19	5.10 +/- 5.67	0.15 +/- 0.49	N/A	N/A
QEW (7)					
Prevalence	0.0%	28.6%	12.5%	85.7%	0.0%
Mean intensity	N/A	0.86 +/- 1.57	0.71 +/- 1.89	N/A	N/A
AP (17)					
Prevalence	0.0%	0.0%	35.3%	47.1	41.2%
Mean intensity	N/A	N/A	0.41 +/- 0.62	N/A	N/A

**Table 1.5.** AICc models with lowest scores ( $\Delta$  AICc < 3) indicating what variables are affecting hemoglobin levels in deer mice (*Peromyscus maniculatus*). IS = blacklegged ticks (*Ixodes scapularis*) abundance, DV = wood ticks (*Dermacentor variabilis*) abundance, and DV Nymphs = wood tick nymph presence.

Model	AICc	$\Delta$ AICc	AICc Wt	Cum. Wt	LL
DV + IS + DV Nymphs	161.97	0.00	0.29	0.29	-75.20
DV + IS + DV:IS + DV Nymphs	164.43	2.46	0.09	0.59	-75.08

**Table 1.6.** Top two models based on AICc scores for what variables affect hemoglobin levels in deer mice (*Peromyscus maniculatus*).

Bolded rows indicate variables where  $p < 0.05$ . AICc scores for each model are in parentheses. IS = blacklegged ticks (*Ixodes scapularis*) abundance, DV = wood ticks (*Dermacentor variabilis*) abundance, and DV Nymphs = wood nymph presence.

Model	Estimate	Std. Error	t - value	Pr(> t )	R <sup>2</sup>
DV + IS + DV Nymphs (161.97)	Intercept: 16.60 DV: -0.25 IS: -0.88 DV Nymphs: 1.95	Intercept: 0.26 DV: 0.06 IS: 0.23 DV Nymphs: 0.77	Intercept: 63.87 DV: -4.01 IS: -3.90 DV Nymphs: 2.55	Intercept: $< 2e^{-16}$ <b>DV: 0.0003</b> <b>IS: 0.0004</b> <b>DV Nymphs: 0.01</b>	Multiple: 0.471 Adjusted: 0.431
DV + IS + DV:IS + DV Nymphs (164.43)	Intercept: 16.63 DV: -0.28 IS: -0.96 DV:IS: 0.04 DV Nymphs: 1.85	Intercept: 0.27 DV: 0.09 IS: 0.29 DV:IS: 0.08 DV Nymphs: 0.80	Intercept: 61.40 DV: -2.97 IS: -3.33 DV:IS: 0.46 DV Nymphs: 2.30	Intercept: $< 2e^{-16}$ <b>DV: 0.005</b> <b>IS: 0.002</b> DV:IS: 0.65 <b>DV Nymphs: 0.03</b>	Multiple: 0.474 Adjusted: 0.420

**Table 1.7.** Pearson correlation coefficient values between predictor variables and hemoglobin levels in deer mice (*Peromyscus maniculatus*). The variable ‘Tick abundance’ includes the abundance of both tick species together, in order to compare correlation variation when blacklegged ticks (*Ixodes scapularis*; IS) and wood ticks (*Dermacentor variabilis*; DV) are interacting together.

Predictor variable	Pearson correlation coefficient (r)
Tick abundance	-0.49
IS abundance	-0.51
DV abundance	-0.50
DV Nymph presence	-0.21

**Table 1.8.** Mean hemoglobin levels from each site including all deer mice (*Peromyscus maniculatus*) hosts, only hosts infested with ticks, and only hosts not infested with ticks. P values given are one-way ANOVA results between hemoglobin levels of deer mice infested with ticks versus mice without ticks. Bolded values indicate variables where  $p < 0.05$ . LP = Long Point Provincial Park, QEW = Queen Elizabeth II Wildlands Provincial Park, AP = Algonquin Provincial Park.

Sites	Mean Hgb levels (g/dL)	Mean Hgb levels with tick infested mice only (g/dL)	Mean Hgb levels with non-infested mice only (g/dL)	Pr(>F)
All Sites	15.78 +/-1.86	15.17 +/-2.00	16.46 +/-1.46	<b>0.012</b>
LP	14.89 +/-1.97	14.89 +/-1.97	N/A	N/A
QEW	16.49 +/- 1.33	17.1 +/-1.41	16.24 +/-1.38	0.492
AP	16.55 +/-1.47	16.90*	16.53 +/-1.52	0.814

\* only one tick present at this site; not a focal species

**Chapter II:**

**Ectoparasite community**

**assemblages found on**

*Peromyscus maniculatus*

**across varying degrees of**

**ixodid tick exposure**

## **Abstract:**

Ectoparasites are fundamental to ecosystems, playing a key role in trophic regulation. Fleas, mites, and ticks are common hematophagous ectoparasites that infest shared mammalian hosts. One common host in Ontario, Canada, is the deer mouse (*Peromyscus maniculatus*). As the climate warms and the geographic ranges of blacklegged ticks (*Ixodes scapularis*) and wood ticks (*Dermacentor variabilis*) expand, their introduction to new ecosystems may alter current ectoparasite communities. At three different sites where exposure to ticks varied (both in terms of tick diversity and abundance), ectoparasite community structures found on deer mouse hosts were compared and analyzed, focusing on species co-occurrences and habitat partitioning on the host. I predicted that when varying tick species were prevalent, ticks would dominate the micro-habitat niches often inhabited by other parasites, thereby significantly altering parasite community structure. My results suggest that blacklegged ticks and wood ticks could have a positive association with each other, but a negative or random association with other ectoparasite species, even when they do not occupy the same micro-habitat niche. Sampling site played a significant role in community assemblages as well, possibly due to the differences in tick exposure. As the ticks' ranges expand and they become more abundant, it is important to understand how their prevalence can potentially alter the dynamics in an ectoparasite community, affecting the transmission of pathogens that may spread within an ecosystem, from one host to another.

**Key Words:** Community ecology, fleas, mites, parasite-host interactions, *Peromyscus maniculatus*, ticks

## Introduction

Parasite species rarely occupy a host in isolation (Hellard *et al.*, 2015). Rather, assemblages of parasites infest an individual host (Bordes *et al.*, 2007; Kappeler *et al.*, 2015). Parasites experience intraspecific and interspecific relationships and directly or indirectly compete with each other (Bordes & Morand, 2011). Interspecific interactions may be synergistic, where one parasite species facilitates the other, or antagonistic, where one species suppresses the other (Fenton *et al.*, 2010; Hellard *et al.*, 2015). For instance, hard ticks (Acari: Ixodidae) can promote infestation rates of other tick species (Lutermann *et al.*, 2015), and parasitic mites (Acari: Mesostigmata) have been found to act as biological controls to mosquito infestations (Atwa *et al.*, 2017). Fleas (Insecta: Siphonaptera) and mites have also been found to experience antagonistic relationships with ticks that they share a host with (Krasnov *et al.*, 2010b; Lutermann *et al.*, 2015; Hoffman *et al.*, 2016).

Not all co-occurring species may have a clearly defined synergistic or antagonistic relationship, however. In one study analyzing the concurrent feeding of two tick species (blacklegged tick, *Ixodes scapularis* and winter tick, *Dermacentor albipictus*) on white-tailed deer (*Odocoileus virginianus*), blacklegged ticks were more abundant and prevalent in general, although their prevalence did not alter the distribution of winter ticks (Baer-Lehman *et al.*, 2012). Blacklegged ticks can be the dominant species on a host when present, although other ectoparasites may be more competitive as blacklegged ticks tend to have lower success rates of attaching to a host (Baer-Lehman *et al.*, 2012). Since it is rare for only one ectoparasitic species to be infesting a host (Bordes & Morand, 2011), and there is a possibility of interspecific competition occurring between ectoparasites (Baer-Lehman *et al.*, 2012), co-occurrences can

affect parasite species diversity on a host, which in turn can cause cumulative effects of parasitism on the ecology and evolution of the host (Bordes & Morand, 2011).

Differences in ecology between parasites may affect interactions between parasitic species sharing a host. Fleas migrate between their host and the host's nest, which can affect their interspecific relationships with other ectoparasites as they are not consistently sharing the host (Krasnov *et al.*, 2010a). Energy requirements for immune responses to parasites can also differ depending on the host species. In one study where two gerbil species (*Gerbillus andersoni* and *Gerbillus dasyurus*) were exposed to fleas, one species was significantly affected by parasitizing fleas more so than the other, suggesting that even closely related organisms may not have similar immune responses to certain parasites (Hawlana *et al.*, 2006).

Host associations within an ecosystem and seasonality differences in macro-habitats (*i.e.*, habitats with complex flora and fauna; landscape ecosystems) may also influence variation in host preferences by parasites (Bouchard *et al.*, 2011). This may result in differences in parasite loads between conspecific hosts from different habitats. For instance, Halsey and Miller (2018) explored host-tick associations between blacklegged ticks and two of their hosts – white-footed mice (*Peromyscus leucopus*) and white-tailed deer – over time. Their results suggest that when parasites and hosts co-evolve, the hosts are able to adapt a more effective grooming technique. At the same time, an adaptive behavioural strategy for ectoparasites to combat host grooming is to feed on areas where it is difficult to groom (Fracasso *et al.*, 2019). Many ectoparasites can feed from several different micro-habitat niches on their hosts (Balashov, 1972), however, depending on the parasite's size and mobility, as well as the host's defense mechanisms, parasites tend to prefer specific parts of the host's body and will partition the micro-habitat with their competitors (Poulin, 2011).

The purpose of this study is to compare the community assemblage of ectoparasites on deer mice from three localities across Ontario, examining the differences between communities with and without ticks. Specifically, I compared ectoparasite attachment sites on hosts, species distributions between sampling landscape sites, and co-occurrences of ectoparasites with varying tick prevalence. I predicted that when ticks were occurring on a host, these ectoparasites would dominate the micro-habitat niches often inhabited by other parasites. There has been evidence that blacklegged ticks can be the more abundant species on a host relative to another tick species, but the impact of tick prevalence in general on the co-occurrence patterns of other ectoparasites is currently unknown (Baer-Lehman *et al.*, 2012). I predicted that when ticks are occurring on an individual host, there would be a negative co-occurrence with other ectoparasites, particularly in the preferred areas on the mouse's body where several species are known to congregate (*i.e.* the ears, where ticks and mites are commonly found; Ostfeld *et al.*, 1993; Baer-Lehman *et al.*, 2012; Bobbie *et al.*, 2016) due to interspecific competition. Knowing how the ectoparasite community changes when ticks are a part of the community is important for understanding how the mouse host will be influenced by an encounter with a newly introduced parasite that can affect parasite diversity as well as the host's ecology and evolution (Bordes & Morand, 2011).

## **Methods**

All methods in this study were approved by the Animal Care Committee (ACC) at Laurentian University, file number 6017269.

## *Field Methods*

Three sites were visited for data and specimen collection from May-August 2019. The sites chosen for this experiment took into consideration the current and predicted ranges of blacklegged ticks and wood ticks in Ontario, Canada (Hamer *et al.*, 2014; Clow *et al.*, 2016; 2017; Minigan *et al.*, 2018), and were areas within the geographical habitat range of deer mice (Bedford and Hoekstra, 2015). These sites were Long Point Provincial Park (LP; 42.5817° N, -80.3952° W), Queen Elizabeth II Wildlands Provincial Park (QEW; 44.7534° N, -78.7844° W) and Algonquin Provincial Park (AP; 45.3402° N, -78.2618° W). At LP, populations of blacklegged ticks and wood ticks are well established (Watson & Anderson, 1976; Lindsay *et al.*, 1991). Only wood ticks are known to be established at QEW (Minigan *et al.*, 2018; P. Careless, 2019, pers. comm.), and neither blacklegged ticks or wood ticks have established in or colonized AP (Hamer *et al.*, 2014; Clow *et al.*, 2016; 2017; Minigan *et al.*, 2018; Public Health Ontario, 2019).

Traplines were set up at QEW and LP, modelled after the long-term traplines used in AP (Fryxell *et al.* 1998; Falls *et al.*, 2007). To replicate these lines, three traplines at least 0.3 km apart from each other were set at the field sites, with twenty Longworth traps placed (Penlon Ltd., Oxford, U.K) per line. Each trapline was 90 m in length and had two Longworth traps 10 m apart from each other every 10 m (Fryxell *et al.*, 1998; Falls *et al.*, 2007). Traps were placed in similar forest types across sites to maintain a consistent or similar habitat, although sites were from different ecozones (Crins *et al.*, 2009). Three of the long-term traplines at AP were exclusively used for this project during this time as well. Traps were baited with water-soaked sunflower seeds and were set half an hour before sunset (2000h-2100h) and checked half an hour before sunrise (0430h-0530h). The traps were checked five days each week; three days

consecutively, with a one-day break, and two more consecutive days to reduce the amount of stress in the mice from repeated captivity or trap response behaviour (Nichols *et al.*, 1984). To account for change in seasonality, and the difference in parasite assemblages that occurs within the spring-summer months, I alternated between these three sites in ascending latitude, spending one week at each site, three times, for a total of three weeks per site.

Deer mice were distinguished from white-footed mice by morphological characteristics: differences in ear length (in mm; Kamler *et al.*, 1998), tail bi-colouration, tail “penciling” (J. Bowman, 2019, pers. comm.), and degree of dorsoventral colouration (Buchholz & Dick, 2017). Deer mice were collected from traps at each site, where the right leg of each mouse was shaved to denote individuals as recaptures. Recaptures were not included in any statistical analysis to avoid bias. Individual mice were examined for 60 seconds for sampling of ectoparasites (Patterson *et al.*, 2013). Individual arthropod specimens were combed off the host using an ethanol sterilized louse comb (Hawlena *et al.*, 2006; Patterson *et al.*, 2013) or plucked off using sterilized tweezers (Bobbie *et al.*, 2016) and placed in a collection vial, preserved in 80% ethanol (Krogmann & Holstein, 2010). During this time, the position on the mouse (ears, head, middle, back, and extremities) where parasites were found was recorded. Extremities included forelimbs, hindlimbs, genitals, and tail. Tick identifications were done by hand using several guides (Lindquist *et al.*, 2016; Dubie *et al.*, 2017), and were later confirmed by Dr. Robbin Lindsay from the National Microbiology Laboratory (Winnipeg, Manitoba, Canada). Fleas were identified using an identification key (Holland, 1985), and later confirmed by Dr. Terry Galloway (University of Manitoba). Dr. Heather Proctor from the University of Alberta was able to identify two taxonomically unique mite groups from the Trombiculidae family, one of which she identified to the genus; *Neotrombicula*.

Age of mice was determined by body mass; mice that were 15 g or less were considered juveniles (Banfield, 1974; Schmidt *et al.*, 2019). Sex and reproductive status (non-reproductive/reproductive) were determined visually. Enlarged testes in males and perforated vagina and presence of nipples in females indicated reproductive individuals (Gaitan & Millien, 2016). No pregnant females were included in this study. Other external factors were also documented, such as field sites where samples were collected, and the temporal session when the samples were collected. This was divided by early (May-June), mid (June-July), and late (July-August) summer of 2019.

### *Statistical Methods*

Measurements of prevalence and intensity of ectoparasites were completed using QPweb (Reiczigel *et al.*, 2019). For the purpose of this study, mean intensity refers to the mean number of ectoparasites found on mice across the population, prevalence refers to the percentage of hosts that have the specified parasite infesting them in a population, and abundance refers to the number of ectoparasites found on individuals (Bush *et al.*, 1997). Statistical analyses were conducted in the R environment (R core Team 2019 [v3.6.1]). For all statistical analyses, the ectoparasite species and groups included were: *I. scapularis*, *D. variabilis*, *O. leucopus*, *Neotrombicula sp.*, and Trombiculidae *sp.* Although in AP, the squirrel tick (*Ixodes angustus*) was also present, there was only one instance of this species present on a single host (n = 1). Thus, this species and its host were not included in any ordination or co-occurrence analyses.

Constrained and unconstrained ordination methods were used to determine relationships between ectoparasites using the ‘vegan’ package (Oksanen *et al.*, 2019 [v3.6.1]). Specifically, correspondence analyses (CA) were performed using presence-absence data of ectoparasites on

hosts. An unconstrained CA was conducted comparing the ectoparasite species and where they were found on individual mice micro-habitat sites (head, ears, middle, back, extremities, and unknown). Species and site scores for each axis were recorded where site scores were weighted averages of species scores.

A canonical correspondence analysis (CCA) was also conducted, where the analysis was constrained by environmental factors: host body mass, sex, age, reproductive status, field site, and session. Eigenvalues were extracted from these data (both constrained and unconstrained). A permutation test for constrained correspondence analysis (permutations = 9999) was performed to test the specified environmental factors for significance to ectoparasite species distributions (Oksanen, 2019). This was done using a significance test for each variable as a ‘marginal’ term in the model such that separate significance tests for each variable would be performed, rather than assessing environmental constraints sequentially, where the order of terms can influence significance (Legendre *et al.*, 2011). A similar test was conducted to determine significance by CCA axes as well, where permutations = 9999 (Legendre *et al.*, 2011; Oksanen, 2019). A non-metric multidimensional scaling (NMDS) model was also performed on parasite loads across individual hosts as well to compare dissimilarity of ectoparasite communities between field site locations (LP, QEW, AP). NDMS is a dissimilarity index where ordination spacing represents distance in a dissimilarity index (Oksanen, 2019). For this analysis, each host was considered a unique site to determine if there were similar species found co-occurring across sampling field sites.

A species pair-wise co-occurrence analysis was also conducted using the package ‘cooccur’ (Griffith *et al.*, 2016 [v3.1]) to determine if co-occurrences between species were positive, negative, or random across micro-habitats (individual hosts) based on a statistical

probabilistic co-occurrence model. Here, each mouse was considered a site, where there was no expectation that parasites should be found at the same macro-habitat location (field site). This R implementation is metric-free, distribution-free, and randomization-free (Griffith *et al.*, 2016) and has a similar methodology as a matrix-level approach (Arita, 2016). In this package, the algorithm calculates the observed and expected frequencies of co-occurrences between each pair of species based on presence/absence data distributed among a set of sampling sites (*i.e.*, deer mouse hosts; Veech, 2013; Griffith *et al.*, 2016). The analysis reports significant positive or negative co-occurrences as well as random co-occurrences based on differences between observed and calculated expected co-occurrences across sites (Griffith *et al.*, 2016; Adair *et al.*, 2018). Expected co-occurrence ( $Q_{\text{exp}}$ ) is measured, such that:

$$Q_{\text{exp}} = \sum(P_j \times j)$$

where  $P_j$  is the probability of co-occurrence at exactly  $j$  sites, and  $j$  is the number of co-occurrence sites (Veech, 2013). When co-occurrences are extremely low or extremely high relative to the expected co-occurrence, these associations are considered to be negatively or positively co-occurring, respectively (Veech, 2013; Griffith *et al.*, 2016). When each host is regarded as a micro-habitat, this analysis does not take into consideration the population source (macro-habitat; LP, QEW, AP) that the host is from. Further analyses were conducted within each population source as well to constrain the results by field site for comparison.

## Results

A sample of 70 individual deer mice hosts across field sites were included in analyses. This included 23 female mice and 47 male mice (Table 2.1). Dr. Terry Galloway identified one flea species from the collected samples across sites: *Orchopeas leucopus*. Dr. Heather Proctor

was able to identify two taxonomically separate groups from the mite specimens collected. One group she identified as the genus *Neotrombicula*. The other group could not be identified beyond family; Trombiculidae. For the purpose of this study, the mite groups will be defined as '*Neotrombicula sp.*' and '*Trombiculidae sp.*', respectively. Across hosts, 162 wood ticks and 39 blacklegged ticks were collected, none of which were adults, along with 22 *O. leucopus*. There were 26 instances of *Trombiculidae sp.* and 10 of *Neotrombicula sp.* present on deer mice (Table 2.2). It should be noted that each field site had a different assemblage of parasites. *Trombiculidae sp.* was not present at LP, *Neotrombicula sp.* and blacklegged ticks were not present at QEW, and neither blacklegged ticks nor wood ticks were present at AP. *O. leucopus* was present at all sites (Table 2.2).

An unconstrained correspondence analysis plot of ectoparasite prevalence based on their location on deer mice showed that both tick species and mite groups aggregated at the ears of their hosts, while *O. leucopus* tended to be found across extremities and back (Figure 2.1). Eigenvalues in the unconstrained model (CA) indicate that axis CA1 explained approximately 74% of the variation within the dataset, with *O. leucopus* being highly differentiated from the other species, having the lowest value and only negative axis CA1 value (-2.240). *O. leucopus*' preferred micro-habitats (back and extremities) also had the most differentiated site scores on axis CA1 (-2.338 and -1.953, respectively; Table 2.3). A NMDS model was performed to determine the species distributions of ectoparasite groups or species across field sites. Blacklegged ticks were only prevalent in LP, although LP presented a somewhat similar species distribution with QEW (Figure 2.2; see the overlapping ellipses) as wood ticks were prevalent at both sites (Figure 2.2; Table 2.2). However, there was greater overlap between the parasite distribution of QEW and AP (Figure 2.2), suggesting a larger overlap in ectoparasite

communities, as both sites have similar prevalence of *O. leucopus* and Trombiculidae *sp.* (Table 2.2). The greatest disparity in species distributions was between LP and AP (Table 2.2).

When comparing eigenvalues of constrained and unconstrained models, the unconstrained model had a larger proportion rank (CA: 0.624, CCA: 0.376) and inertia (CA: 1.259, CCA: 0.759), however, the axis CCA1 in the constrained model explains the greatest proportion of variance overall in the constrained model (0.333; Table 2.4). On this axis, field site had the highest biplot score for constraining variables (0.991) and both tick species had the lowest species scores (*I. scapularis*: -0.608, *D. variabilis*: -0.400; Table 2.5). A permutation test examining the importance of constraining environmental variables assessed that field site significantly affected the ectoparasite communities across hosts ( $p = 0.0001$ ; Table 2.6). A permutation test determined that axis CCA1 was the most significant axis in this model as well ( $p = 0.0001$ ; Table 2.6).

My analysis of relationships among ectoparasites detected 10 pairwise associations of statistically significant species co-occurrences when considering hosts as unique habitats/sites. The only positively co-occurring pair was blacklegged ticks and wood ticks (Figure 2.3), which were expected to occur across only 13.7 sites, but were found together at 22 (Table 2.7). Both tick species had negative associations with the mite groups and *O. leucopus* (Figure 2.3), where the expected co-occurrences were much lower than the observed co-occurrences (Table 2.7). The co-occurrences between the mite groups and *O. leucopus*, however, appeared to be random (Figure 2.3), such that species pairs were not observed more or less than expected by random chance. All co-occurrence analyses constrained by field sites determined that the co-occurrences between all ectoparasites were random, suggesting none of these associations between ectoparasites were positively or negatively co-occurring at LP, QEW, or AP (Table 2.8).

## Discussion

### *Ectoparasite Distributions Across Field Sites*

The purpose of this study was to compare community assemblage of ectoparasites on deer mice from populations with varying tick exposure. I hypothesized that when ticks were infesting a host, they would dominate the micro-habitat so that few or no other ectoparasite species (fleas and mites) would be sharing the host, especially in preferred attachment sites (*i.e.* ears). Although studies have found that host parameters such as body mass and sex affect parasite community assemblages (Gaitan & Millien, 2016; Sponchiado *et al.*, 2016), this was not the case in my study. However, field site played an important role in the observed structure. The three field sites that were assessed are geographically different, occurring at different latitudes and in different ecozones (Crins *et al.*, 2009). LP is the most southern site and was the only field site where blacklegged ticks were prevalent. Although wood ticks were prevalent at LP and QEW, there were greater ectoparasite community similarities between QEW and AP. At these two sites, *O. leucopus* and Trombiculidae *sp.* were more prominent than the other ectoparasite species. The intensity of wood ticks in QEW was low and did not seem to affect the occurrence of other ectoparasites found at this site. There are several possible reasons why this may be when comparing QEW and LP, where wood ticks were present: 1) the occurrence of blacklegged ticks affects the prevalence of other ectoparasites more than wood ticks, 2) the intensity of wood ticks affects the infestation rates of co-occurring ectoparasites more so than its presence on a host does, 3) the field site location is affecting the species distribution in a way that was not measured (*i.e.* abiotic factors such as weather or temperature, differences in flora or fauna between habitats). Field site had the highest biplot score for constraining variables, significantly affecting the ectoparasite communities across hosts, and it is possible that the abiotic environmental

factors of these sites are influencing ectoparasite communities (Gómez-Rodríguez *et al.*, 2015). Thus, there may be environmental characteristics not considered in this study that are influencing occurrences of different ectoparasites.

### *Ectoparasite Distributions Across Hosts*

The correspondence analysis plot illustrates that both tick species and both mite groups prefer the ears as their attachment site, although wood ticks can also be found on the head and middle of their host's body (Figure 2.1). Fleas, on the other hand, were never found on the head of any of the deer mice, suggesting some exclusion or difference in preference in attachment or feeding sites. Siphonaptera (flea) species differ from Acari species (mites and ticks) in that they invest more time associating with their host, alternating between the host's body and the host's nest (Krasnov *et al.*, 2010a). While all three groups are obligate parasites, fleas do not attach to one site on the host to feed nor do they move on to another host as ticks do (Krasnov *et al.*, 2010a), which can affect associations seen in this study.

Despite fleas moving freely across their hosts, the micro-habitat partitioning of mites and ticks infesting deer mice in this analysis was similar to the body partitioning results found in a study on the painted spiny pocket mouse (*Liomys pictus*) in Mexico (Gómez-Rodríguez *et al.*, 2015). Although the painted spiny pocket mouse lives in an area with generally higher biodiversity than the deer mice in this study, both host species were observed to have ectoparasite communities with mutually exclusive areas on the body occupied by certain ectoparasite groups. On painted spiny pocket mice, hard tick species, including *Ixodes* sp. were found mainly on the head (neck, cheek pouch, and snout), and while some mite species were found exclusively on the ears of their hosts, others were found across the body, from the dorsal

posterior to the ears (Gómez-Rodríguez *et al.*, 2015). The mite species in this thesis were also unknown; yet, two distinct groups were defined: *Neotrombicula sp.* was found in LP and AP and Trombiculidae *sp.* in AP and QEW. *Neotrombicula sp.* is known to infest their host's ears in AP (Bobbie *et al.*, 2016), and were found to be prevalent at this field site. Although also prevalent in LP, they were only found on two individual hosts, while tick prevalence was extremely high at this site. Thus, it may be possible that mutual exclusion of micro-habitats is occurring here, where the tick species are outcompeting the mites that prefer the same attachment sites. It is also possible that at this site, *Neotrombicula sp.* associates with a different rodent host and so their prevalence was not as high in this area. Without knowing the actual species of mite, however, it is difficult to discern its ecology.

#### *Ectoparasite Co-occurrences*

Results differed in co-occurrences when considering hosts as unique micro-habitats versus hosts constrained by their macro-habitats (field site/population source). When hosts were considered individual micro-habitats blacklegged ticks and wood ticks were positively co-occurring, and both tick species were negatively co-occurring with Trombiculidae *sp.* and fleas. Wood ticks were also negatively co-occurring with *Neotrombicula sp.*, whereas blacklegged ticks and *Neotrombicula sp.* were randomly co-occurring. All other co-occurrences were random (Figure 2.3).

It is common for different tick species to feed from the same host (Baer-Lehman *et al.*, 2012; Gómez-Rodríguez *et al.*, 2015; Cayol *et al.*, 2018). In some cases, the incidence rate of a tick species will significantly increase when their host is co-infested by other tick species (Biguezoton *et al.*, 2016). There are several cases where an ectoparasite species infesting a

mammalian host will have increased intensities when there are co-infestations by other taxa on the host, often due to how these interactions compromise the host's immune response (Matthee & Krasnov, 2009; Krasnov *et al.*, 2010b; Lutermann *et al.*, 2015). It is also possible for one tick species to facilitate another tick species in a parasite community, however; it is typical for these interspecific relationships to be antagonistic (Matthee & Krasnov, 2009; Lutermann *et al.*, 2015). A study on eastern rock elephant shrew (*Elephantulus myurus*) ectoparasite communities reported that when experimentally reducing abundances of several tick and flea species from hosts, most inter-taxon relationships of ectoparasites were antagonistic, although some tick species interactions resulted in facilitation of one another (Lutermann *et al.*, 2015). Much like Lutermann *et al.*'s (2015) results, the positive co-occurrence between blacklegged ticks and wood ticks suggests that the competition for attachment sites or blood-feeding resources does not dictate the broad co-occurrence patterns between these species.

While mite and flea co-occurrences were random, there were marked negative co-occurrences found between blacklegged ticks and Trombiculidae *sp.*, and *O. leucopus* as well as between wood ticks and Trombiculidae *sp.*, *Neotrombicula sp.*, and *O. leucopus*. An experimental study that removed a dominant hard tick species from hosts found that the removal of this ectoparasite increased the abundances of co-infesting chiggers and lice (Hoffman *et al.*, 2016). Similarly, in my study, when blacklegged ticks and wood ticks were not present on a host, the prevalence of *O. leucopus* and mites was higher in deer mice. These results are also similar to Lutermann *et al.*'s (2015) findings, where fleas were more prevalent when ticks were removed and vice versa. This may be due to direct competition for resources. Although fleas and mites parasitize *Peromyscus* mice for long periods of time and are consistently impacting their host's immune response (Mize *et al.*, 2011), ticks require larger blood meals for shorter periods of time

(McKenzie, 1998), and since they require greater volumes of a shared resource, they may be outcompeting other blood-feeding ectoparasites. Unfortunately, since surveying only occurred over a short field season between May and August of one year – and the hosts' nests were not examined – sampling effort was limited, so there could be hidden effects impacting both the host and parasite fitness that could be noted over a longer time period (Krasnov & Matthee, 2010).

When micro-habitats were constrained by their population sources, all co-occurrences between ectoparasites were considered random. This result suggests that there are no associations between any of the ectoparasites, though the unbalanced and small sample sizes for each population source could be affecting the results, as expected pair-wise co-occurrences are the product of the two species' probability of occurrence multiplied by the number of sampling sites (Veech, 2013; Griffith *et al.*, 2018). As mentioned previously, sampling effort could also be affecting these results. Although it is possible for co-occurrences between species to be random, it seems unlikely that all species in this study are randomly co-occurring, since many of these ectoparasites favoured the same micro-habitat niches (specifically the ears). In order to further test co-occurrences between these species, an experimental study in which a parasite is removed from the micro-habitat and the parasite community is re-examined (Lutermann *et al.*, 2015) should be conducted.

### *Conclusion*

This study focused on parasite community assemblage processes at the host individual level. However, both population and (macro) community dynamics can affect an individual's parasite load and must also be considered (Gómez-Rodríguez *et al.*, 2015; Hellard *et al.*, 2015). Specifically, interactions at one ecological level can cascade up to other levels, affecting

intraspecific and interspecific relationships between organisms (Hellard *et al.*, 2015). It is especially important to consider these other interactions, as several pathogenic parasites are expanding their geographic ranges northwards (Gasami *et al.*, 2018), which may alter current associations between species. The results of my study suggest that the prevalence of blacklegged ticks and wood ticks could affect community assemblages. As these species expand their ranges (Leighton *et al.*, 2012; Gasami *et al.*, 2018; Minigan *et al.*, 2018), it is possible that they will outcompete other ectoparasites in their micro-habitat.

There is evidence that tick-borne diseases do expand with the increased distribution of *Ixodes* ticks and it is likely that this trend will continue (Ostfeld & Brunner, 2015). Many of the ectoparasites currently found on deer mice (*Peromyscus maniculatus*) are vectors for several pathogens that can be transmitted to humans (Lindsay *et al.*, 2001; DeNatale *et al.*, 2002; Westblade *et al.*, 2017; Mlera *et al.*, 2018; Talagrand-Reboul *et al.*, 2018), indicating the epidemiological importance of these hosts. Since the blacklegged tick and wood tick are vectors of pathogens that can be transmitted to humans (blacklegged tick: *Borrelia burgdorferi*; Sonenshine, 2018; wood tick: *Rickettsia rickettsii*; Durden *et al.*, 2004) and these tick species are expanding their geographical ranges throughout Ontario (Leighton *et al.*, 2012; Clow *et al.*, 2016; 2017; Minigan *et al.*, 2018), it is an important next step to see how they are associating with other ectoparasites that may also transmit zoonotic pathogens.

By understanding the differences in community assemblages between host populations where ticks are present or absent, we can better predict how ectoparasite communities may change when these tick species do eventually expand northwards into new populations. Furthermore, more research is needed to better understand how their presence may affect

parasite-parasite interactions as well as the transmission of zoonotic pathogens that may be able to spread between parasites and hosts.

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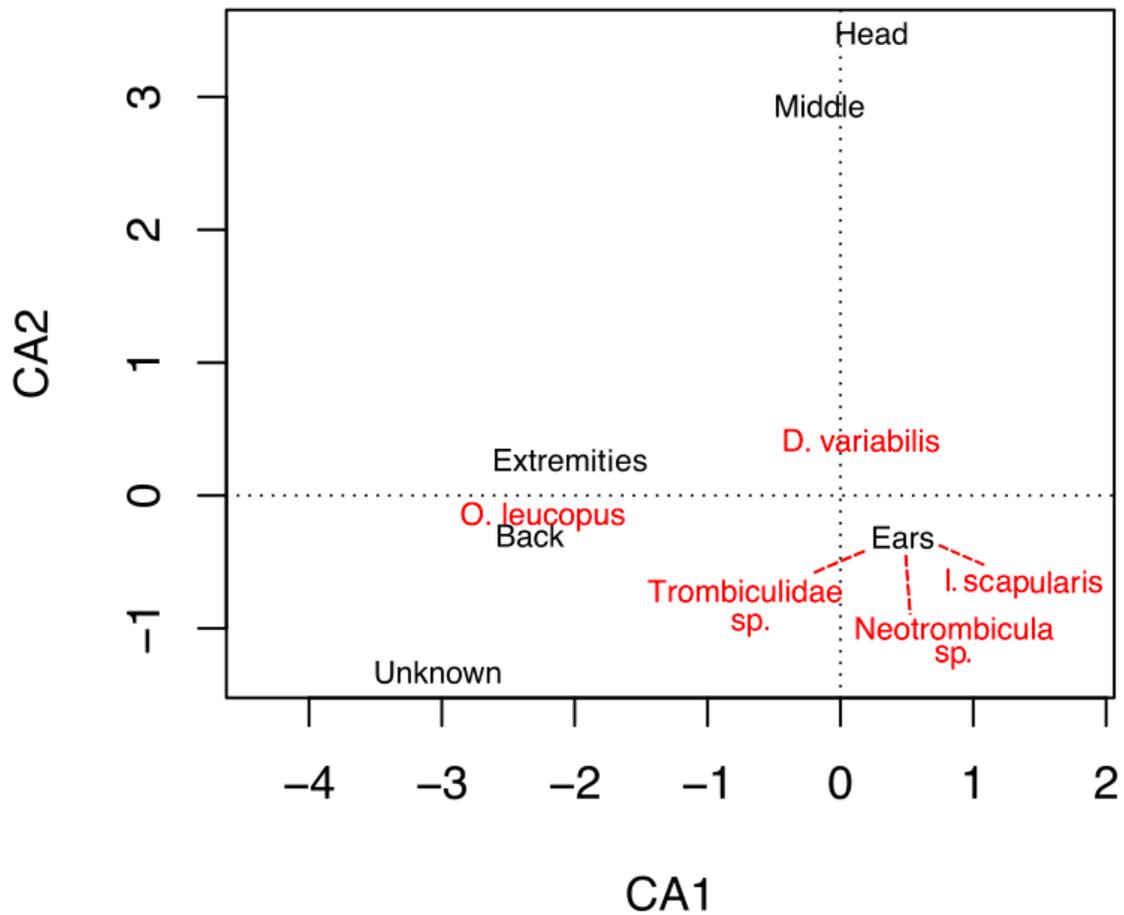
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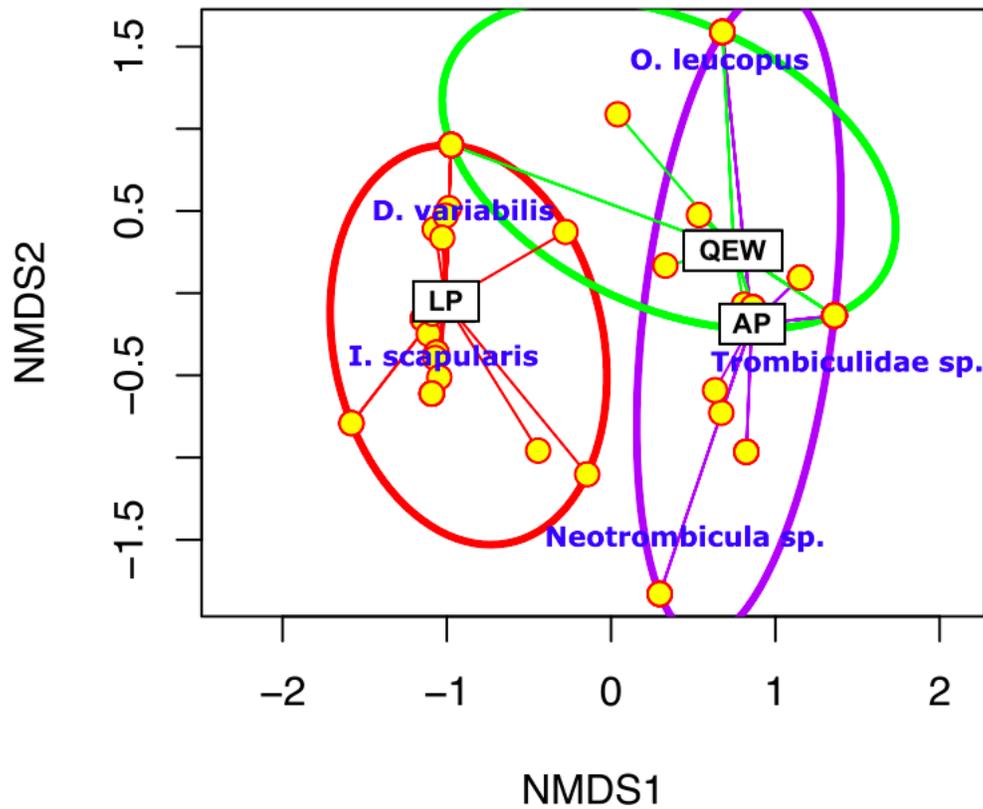
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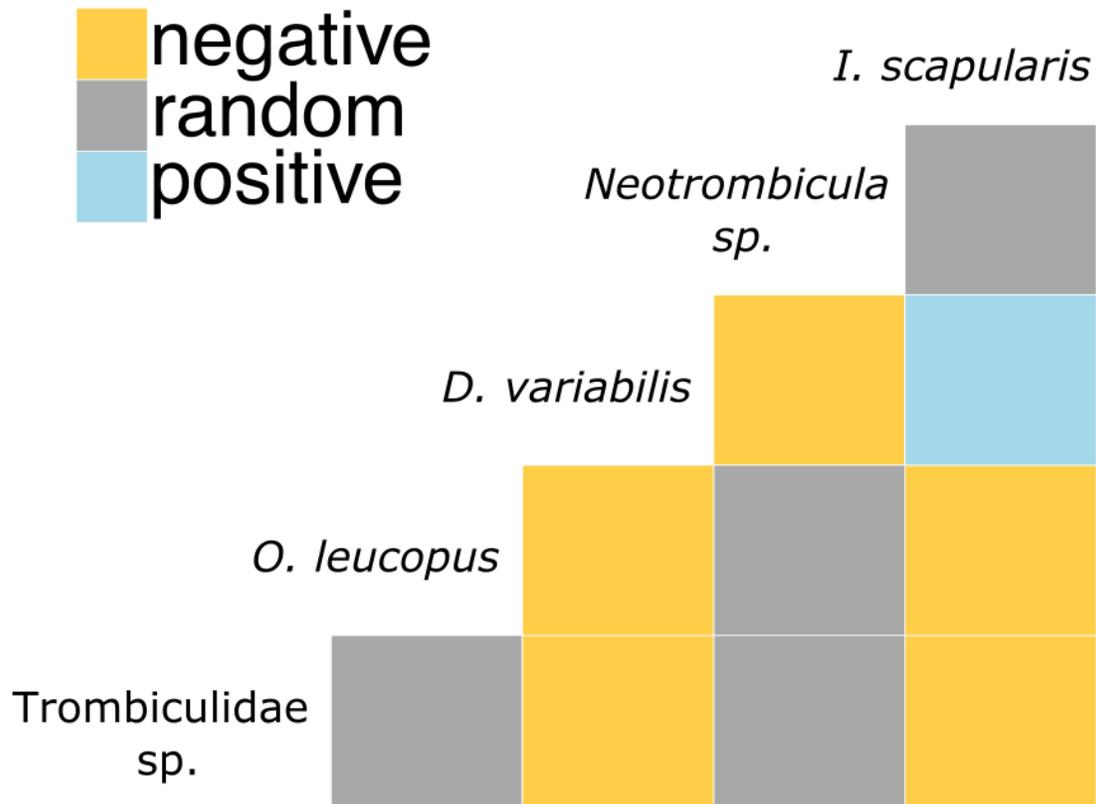
## Figures and Tables



**Figure 2.1.** Correspondence analysis plot of ectoparasite (red text) prevalence to body location (black text) on deer mice (*Peromyscus maniculatus*; n = 70). Dotted red lines indicate that these groups were overlapping on the specified body location.



**Figure 2.2.** Non-metric multidimensional scaling plot of individual deer mice (*Peromyscus maniculatus*; n = 70) and the ectoparasites they host across field sites. Each yellow dot represents a single host (micro-habitat site). The coloured lines extending out from the site names are measurements of dissimilarity, whereby longer lines indicate greater dissimilarity. Red ellipse represents the distribution of Long Point (LP) individuals (hosts and parasites), green ellipse represents Queen Elizabeth II Wildlands (QEW), and purple ellipse represents Algonquin (AP) individuals.



**Figure 2.3.** Heat map depicting the positive, negative, and random species associations determined by the probabilistic co-occurrence model (Table 2.7) for ectoparasites found across individual deer mice (*Peromyscus maniculatus*) hosts (n = 70). Species names are positioned to indicate the columns and rows that represent their pairwise relationships with other species.

**Table 2.1.** Host parameters for deer mice (*Peromyscus maniculatus*) depicting distributions of average body mass (with standard deviation), sex, age, and reproductive status across all sites and within sites as well as temporal sessions. Numbers in parentheses indicate the sample size of hosts for each indicated category. NR = non reproductive and R = reproductive. LP = Long Point Provincial Park, QEW = Queen Elizabeth II Wildlands Provincial Park, AP = Algonquin Provincial Park.

	All Sites (70)	LP (31)	QEW (20)	AP (19)
Average body mass (g)	19.67 +/- 5.28	20.21 +/- 6.67	18.73 +/- 3.85	19.79 +/- 3.95
Host sex				
	Male (47)	Male (21)	Male (13)	Male (13)
	Female (23)	Female (10)	Female (7)	Female (6)
Host age				
	Adult (59)	Adult (26)	Adult (15)	Adult (18)
	Juvenile (11)	Juvenile (5)	Juvenile (5)	Juvenile (1)
Host reproductive status				
	NR (37)	NR (22)	NR (11)	NR (4)
	R (33)	R (9)	R (9)	R (15)
Session				
	Early (10)	Early (5)	Early (3)	Early (2)
	Mid (25)	Mid (12)	Mid (8)	Mid (5)
	Late (35)	Late (14)	Late (9)	Late (12)

**Table 2.2.** Prevalence and mean intensity of ectoparasites found on deer mice (*Peromyscus maniculatus*) across all sites and per site. Numbers in parentheses indicate the number of hosts (first column) and parasites for each indicated column. It should be noted that both mite groups had prevalence measured only; intensities unknown. LP = Long Point Provincial Park, QEW = Queen Elizabeth II Wildlands Provincial Park, AP = Algonquin Provincial Park.

	<i>I. scapularis</i>	<i>D. variabilis</i>	<i>O. leucopus</i>	Trombiculidae <i>sp.</i>	<i>Neotrombicula</i> <i>sp.</i>
All Sites (70)	(39)	(162)	(22)	(26)	(10)
Prevalence	54.3%	55.6%	30.5%	37.1%	14.3%
Mean intensity	0.56 +/- 0.97	2.33 +/- 3.94	0.31 +/- 0.77	N/A	N/A
LP (31)	(39)	(139)	(2)	(0)	(2)
Prevalence	77.4%	93.5%	3.2%	0.0%	6.5%
Mean intensity	1.25 +/- 1.12	4.52 +/- 4.86	0.06 +/- 0.36	N/A	N/A
QEW (20)	(0)	(23)	(10)	(14)	(0)
Prevalence	0.0%	55.0%	25.0%	70.0	0.0%
Mean intensity	N/A	1.15 +/- 2.08	0.50 +/- 1.19	N/A	N/A
AP (19)	(0)	(0)	(10)	(12)	(8)
Prevalence	0.0%	0.0%	42.9%	63.1	42.1%
Mean intensity	N/A	N/A	0.53 +/- 0.61	N/A	N/A

**Table 2.3.** Species and site scores for unconstrained correspondence analysis (Figure 2.1) of ectoparasite prevalence to micro-habitat sites on deer mice (*Peromyscus maniculatus*) hosts (n = 70). Eigenvalues and proportions explained for both CA axes are also included. Inertia = 0.859 (scaled chi-square).

	CA1	CA2
Eigenvalues	0.739	0.119
Proportion explained	0.861	0.139
Species scores		
<i>D. variabilis</i>	0.148	0.413
<i>I. scapularis</i>	0.469	-0.316
<i>O. leucopus</i>	-2.240	-0.159
Trombiculidae <i>sp.</i>	0.469	-0.316
<i>Neotrombicula sp.</i>	0.469	-0.316
Site scores		
Head	0.299	3.767
Ears	0.469	-0.316
Middle	-0.158	2.930
Back	-2.338	-0.304
Extremities	-1.953	0.267
Unknown	-3.030	-1.331

**Table 2.4.** Correspondence analysis inertia (scaled chi-square), proportion ranks, and eigenvalues for both unconstrained and constrained axes for parasite loads found on individual deer mice (*Peromyscus maniculatus*) hosts (n = 70). Proportion explained for eigenvalue contribution to scaled chi-square is in parentheses. Constrained environmental factors include host body mass, sex, age, reproductive status, field site, and session. Mite groups only provided prevalence data, while both tick species and *O. leucopus* provided intensity values.

	Constrained (CCA)	Unconstrained (CA)
Inertia	0.759	1.259
Proportion Rank	0.376	0.624
Axis 1	0.672	0.457
Eigenvalues	(0.333)	(0.227)
Axis 2	0.070	0.450
Eigenvalues	(0.035)	(0.223)
Axis 3	0.015	0.268
Eigenvalues	(0.007)	(0.133)
Axis 4	0.002	0.084
Eigenvalues	(0.001)	(0.040)

**Table 2.5.** Species scores and biplot scores for constraining variables in constrained (CCA) and unconstrained (CA) correspondence analyses for ectoparasite communities across individual deer mice (*Peromyscus maniculatus*) hosts (n = 70). Since CA is unconstrained, all biplot scores for constraining variables are equal to zero. HRS = host reproductive status.

	CCA1	CCA2	CCA3	CCA4	CA1	CA2
<b>Species scores</b>						
<i>Trombiculidae sp.</i>	1.518	-0.248	-0.091	-0.077	-1.613	0.147
<i>Neotrombicula sp.</i>	1.744	-0.706	0.255	0.119	1.019	-2.839
<i>O. leucopus</i>	1.385	0.673	-0.068	0.044	1.300	1.270
<i>I. scapularis</i>	-0.608	-0.191	-0.239	0.035	0.195	-0.127
<i>D. variabilis</i>	-0.400	0.039	0.069	-0.009	-0.018	0.009
<b>Biplot scores</b>						
Host body mass	-0.106	-0.234	-0.124	0.510	0	0
Host sex	-0.072	-0.290	0.144	-0.810	0	0
Host age	-0.070	-0.482	0.293	0.329	0	0
HRS	0.287	-0.678	-0.499	0.374	0	0
Site	0.991	0.010	-0.002	0.042	0	0
Session	-0.060	-0.571	0.728	0.014	0	0

**Table 2.6.** Permutation test results for constrained correspondence analysis (permutations = 9999). The test was performed to assess the specified environmental factors for significance to ectoparasite communities across deer mice (*Peromyscus maniculatus*) hosts (n = 70). Bolded terms indicate significant alpha values (< 0.05). Permutation test results for CCA axes significance is also included here (permutations = 9999). Bolded terms indicate significant alpha values (< 0.05). HRS = Host reproductive status.

	Df	Chi Square	F	Pr(>F)
<b>Environmental Factors</b>				
Host body mass	1	0.007	0.343	0.840
Host sex	1	0.010	0.524	0.699
Host age	1	0.013	0.641	0.618
HRS	1	0.035	1.773	0.117
Field site	1	0.660	33.029	<b>0.0001</b>
Session	1	0.033	1.644	0.155
Residual	63	1.259	N/A	N/A
<b>Axis Significance</b>				
CCA1	1	0.672	34.691	<b>0.0001</b>
CCA2	1	0.070	3.615	0.333
CCA3	1	0.015	0.774	0.991
CCA4	1	0.002	0.080	1.000
Residual	65	1.259	N/A	N/A

**Table 2.7.** Summary of probabilistic co-occurrence model of pair-wise co-occurrences for ectoparasites found across individual deer mice (*Peromyscus maniculatus*) hosts (n = 70) that were each considered a site (micro-habitat) in this analysis.

Species 1	Species 2	Number of micro-habitats that have species 1	Number of micro-habitats that have species 2	Observed number of micro-habitats having both species	Expected number of micro-habitats having both species	Probability of micro-habitats having both species	Probability of co-occurrence freq. < observed # of co-occurrence micro-habitats if both species were distributed randomly	Probability of co-occurrence freq. > observed frequency
<i>Neotrombicula sp.</i>	Trombiculidae <i>sp.</i>	27	10	5	3.9	0.055	0.875	0.321
<i>Neotrombicula sp.</i>	<i>O. leucopus</i>	27	15	7	5.8	0.083	0.847	0.331
<i>Neotrombicula sp.</i>	<i>I. scapularis</i>	27	24	0	9.3	0.132	0.000	1.000
<i>Neotrombicula sp.</i>	<i>D. variabilis</i>	27	40	8	15.4	0.220	0.001	1.000
Trombiculidae <i>sp.</i>	<i>O. leucopus</i>	10	15	2	2.1	0.031	0.636	0.686
Trombiculidae <i>sp.</i>	<i>I. scapularis</i>	10	24	1	3.4	0.049	0.077	0.990
Trombiculidae <i>sp.</i>	<i>D. variabilis</i>	10	40	2	5.7	0.082	0.013	0.999
<i>O. leucopus</i>	<i>I. scapularis</i>	15	24	1	5.1	0.073	0.009	0.999
<i>O. leucopus</i>	<i>D. variabilis</i>	15	40	4	8.6	0.122	0.008	0.999
<i>I. scapularis</i>	<i>D. variabilis</i>	24	40	22	13.7	0.196	1.000	0.000

**Table 2.8.** Summary of probabilistic co-occurrence model of pair-wise co-occurrences for ectoparasites found across individual deer mice (*Peromyscus maniculatus*) hosts (n = 70) that were each considered a site (micro-habitat) in this analysis, constrained by field site (macro-habitat). Numbers in parentheses indicate the sample size of hosts for each site. QEW = Queen Elizabeth II Wildlands.

Species 1	Species 2	Number of micro-habitats that have species 1	Number of micro-habitats that have species 2	Observed number of micro-habitats having both species	Expected number of micro-habitats having both species	Probability of micro-habitats having both species	Probability of co-occurrence freq. < observed # of co-occurrence micro-habitats if both species randomly distributed	Probability of co-occurrence freq. > observed frequency
Long Point (31)								
<i>Neotrombicula sp.</i>	<i>O. leucopus</i>	2	1	0	0.1	0.002	0.936	1.000
<i>Neotrombicula sp.</i>	<i>I. scapularis</i>	2	24	1	1.5	0.050	0.407	0.955
<i>Neotrombicula sp.</i>	<i>D. variabilis</i>	2	29	2	1.9	0.060	1.000	0.873
<i>O. leucopus</i>	<i>I. scapularis</i>	1	24	1	0.8	0.025	1.000	0.774
<i>O. leucopus</i>	<i>D. variabilis</i>	1	29	1	0/9	0.030	1.000	0.936
<i>I. scapularis</i>	<i>D. variabilis</i>	24	29	22	22.5	0.724	0.594	1.000
QEW (20)								
Trombiculidae sp.	<i>O. leucopus</i>	15	5	2	3.8	0.188	0.073	0.995
Trombiculidae sp.	<i>D. variabilis</i>	15	11	8	8.2	0.413	0.605	0.779
<i>O. leucopus</i>	<i>D. variabilis</i>	5	11	3	2.8	0.138	0.779	0.605
Algonquin (19)								
Trombiculidae sp.	<i>Neotrombicula sp.</i>	12	8	5	5.1	0.266	0.663	0.703
Trombiculidae sp.	<i>O. leucopus</i>	12	9	5	5.7	0.299	0.430	0.870
<i>Neotrombicula sp.</i>	<i>O. leucopus</i>	8	9	2	3.8	0.199	0.115	0.985

## General Discussion

There is a clear difference in the hematology and community assemblage of deer mice when tick exposure varies, particularly when both blacklegged ticks and wood ticks are prevalent. The prevalence of both blacklegged ticks and wood ticks has been demonstrated to affect their host's hematology and ectoparasite community structures. Deer mice that were infested with blacklegged ticks had hemoglobin levels significantly lower than mice that were not infested. In most cases, blacklegged ticks were found on mice with wood ticks. Mice with higher intensities of wood ticks also had significantly lower hemoglobin levels, specifically in LP. LP had the highest tick intensities and was the only site where blacklegged ticks were prevalent. Without confounding effects, these two species affected their host's hematology separately, as their interaction was found to be nonsignificant, although the results in Chapter 2 suggest these species do often occur together. Wood tick nymphs, which are larger than larval ticks (Lindquist *et al.*, 2016), were also shown to have a negative effect on hemoglobin levels.

Low hemoglobin levels can result in iron deficiency and anemia (Andrews, 1997; O'Brien *et al.*, 2003). In lab studies focused on rodents, there is evidence that iron deficiency causes fatigue, trouble focusing, reduced immune response, and loss of motor skills (Rosales *et al.*, 1999; Nicolas *et al.*, 2002; Hentze *et al.*, 2004; Camaschella, 2015). This could in turn affect the range expansion of the infested host (Binning *et al.*, 2017), potentially suppressing the expansion of pathogens they carry to new populations by migrating less (Huang *et al.*, 2019) or intensifying the spread of pathogens within their current one by reducing genetic diversity (King & Lively, 2012). Since other variables, such as the host's microbiota, endoparasite loads, and diseases were not included in any analysis, there are still many questions that must be answered

to fully understand the deer mouse's hematology, although the prevalence of the two tick species appears to affect hemoglobin levels.

These two species of ticks also appeared to affect the ectoparasite community assemblages of their hosts. It is possible that there is a positive co-occurrence association between the two tick species. Similar to the hematology results, blacklegged ticks and wood ticks were often found together and did not seem to deter each other. Again, LP was the only site examined that had both tick species, where most of the hosts had tick infestations, but few had fleas or mites prevalent. In contrast, the greatest prevalence of mites and fleas was seen in AP - where neither tick species has currently established. In the unconstrained analysis, it was found that while blacklegged ticks and wood ticks were positively co-occurring with each other, these ticks negatively co-occurred with fleas and mites (except blacklegged ticks and *Neotrombicula sp.*, which were random), whereas fleas and mites were randomly co-occurring. The absence of co-occurrence between ticks and mites could be due to interspecific competition, as they all preferred the same attachment site/microhabitat niche (ears). However, fleas were also negatively co-occurring with ticks, despite partitioning the habitat, possibly due to differences in their ecology and host reliance (short term versus long term infestations).

Field sampling site also played a significant role in species distributions in my system. When micro-habitats were segregated by their macro-habitats, all co-occurrences between ectoparasites were deemed random. Although the total number of ectoparasite species found across sites was not different, there was a difference in tick species prevalent at each site. Host populations at lower latitudes have been documented to have higher parasite species diversity (Preisser, 2019; Bordes *et al.*, 2011). Although, the latitudinal range in this study was small, each site was in a different ecozone (Crins *et al.*, 2009). Differences in habitat types can have

significant effects on parasite loads (Carbayo *et al.*, 2019) as well as differences in host population sizes (Papkou *et al.*, 2016) so these variables should be considered in further studies. Similar to the host's hematology, an individual's microbiota, endoparasites, and pathogens can also affect their parasite community assemblage (Budischak *et al.*, 2012) and should be examined as well.

The results of my thesis suggest that further consideration must be given on how the ever-increasing prevalence of these tick species will affect ecosystems that will soon come into contact with them. QEW is expected to have blacklegged ticks invade in the upcoming years (Leighton *et al.*, 2012; Ogden *et al.* 2013; Clow *et al.*, 2016; 2017), and soon after AP will be invaded by both tick species (Leighton *et al.*, 2012; Ogden *et al.* 2013; Minigan *et al.*, 2018). As naïve hosts come into contact with these tick species, it is important to understand how tick prevalence will affect the host's physiology via hematology as well as their ectoparasite communities. Deer mice in particular, are vectors to numerous pathogens, passed on to numerous other species via ticks and other ectoparasites (Lindsay *et al.*, 2001; DeNatale *et al.*, 2002; Westblade *et al.*, 2017; Mlera *et al.*, 2018; Talagrand-Reboul *et al.*, 2018). If prevalence or intensity of ticks reduces the prevalence of other ectoparasites found on a host or within a host's population, this will in turn reduce parasite species richness, thereby reducing biodiversity. Although it has been suggested that higher host biodiversity can dilute the spread of tick-borne diseases (Ogden & Tsao, 2009; Dantas-Torres, 2015), parasite diversity may also act as a control for the spread of zoonotic pathogens (Johnson *et al.*, 2013). This is important to consider, as tick-borne viruses are arguably of the greatest medical concern in North America, particularly those transmitted by the blacklegged tick (Kocan *et al.*, 2015; Kulkarni *et al.*, 2015; Eisen & Eisen, 2018). It has been documented that tick burdens do not affect the survival of *Peromyscus* hosts

(Hersh *et al.*, 2014), and that these hosts may have higher innate immune functions that allow them to enhance their tolerance for high tick burdens compared to other rodents (such as *Microtus* spp.; Rynkiewicz *et al.*, 2013). Since the effects of low hemoglobin levels via parasite loads have not been previously tested on wild *Peromyscus*, it is not known how this deficiency may alter their physiology or immunology.

Further studies on interactions between these tick species and *Peromyscus* hosts should be conducted, particularly between experienced hosts and naïve hosts that may respond differently to these parasitic interactions, physically and immunologically. Since these tick species are expanding their ranges into areas where naïve hosts inhabit (Bedford and Hoekstra, 2015; Clow *et al.*, 2016; 2017; Minigan *et al.*, 2018), future research should focus on how these interactions may differ. Understanding these dynamics can allow us to better predict how new host populations will react to ixodid ticks. Other parasite interactions should also be considered, including associations occurring between flea, mite, and lice species, as well as other tick species currently prevalent in these areas and those tick species that are expected to expand their ranges into these regions in the future. Recognizing these parasite-parasite relationships will allow for greater insight into changes in species richness and biodiversity in a micro-habitat as well as provide information on the transmission of diseases that can spread from host to parasite (and vice versa) or even possibly indirectly from one parasite to another.

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