Eco-epidemiology of Anaplasmosis in the peri-domestic environment of southern Quebec



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INTRODUCTION

- Human granulocytic anaplasmosis (HGA) is a vector-borne disease caused by the bacterium Anaplasma phagocytophilum (Ap).
- The zoonotic "human active" variant (Ap-ha) is pathogenic for humans, horses and dogs.
- 2021: Outbreak of HGA reported in the Estrie region (16/25 cases in Bromont).
- Objective : determine which species of wild small mammals can act as competent reservoir hosts for the zoonotic variant Ap-ha.

METHODS

- Eight peri-domestic woodland sites in the city of Bromont (Estrie): four sites close (< 3km) and four sites far (> 3 km) from human cases.
- June to August 2022 and 2023 : capture of small mammals (live traps) and drag sampling.
- Samples : whole-blood and larvae (small mammals), questing nymphs.
- **Diagnosis :** Multiplex real-time PCR and genetic strain identification.

RESULTS (2022)

- 339 small mammals from 9 species.
- Blood*: 18/93 PCR positive rodents (19.4%).
 Ap-ha strain confirmed in 17 individuals.
- Larvae*: 57,1% of eastern chipmunks (4/7), 15,1% of *Peromyscus* mice (11/73) and 11,1% of red squirrels (1/9) transmitted Ap to at least one larva.
- Prevalence of infection for Ap (positive blood and/or larvae sample)*: 66.7% in eastern chipmunks (10/15), 15.5% in *Peromyscus* mice (16/103) and 14,3% in red squirrels (3/21).
 *Animals recaptured between trapping periods were excluded.
- Questing nymphs: 5.3% positive for Ap (22/412).
- Probability of sampling a positive nymph: 4 times higher at sites close to human cases (p=0.0048).

DISCUSSION

- Other rodent species than *Peromyscus* mice should be considered as potential reservoir hosts for the zoonotic strain of anaplasmosis.
- Results will help better understand HGA emergence factors and guide development of risk management interventions.



Eastern chipmunks have the potential to be important reservoir hosts for anaplasmosis





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Prevalence of Ap (%) in blood and/or larvae samples by species



Proportion of the Ap variants (%) in positive questing nymphs (n=22)



Prevalence of Ap in questing nymphs by site category

Site	No.	Ap [%]	Ap-ha	Ap-var	Mean nymph
	tested				density/100m ²
< 3 km	217	18* [8.3]	11	6	1.37
> 3 km	195	4 [2.1]	3	1	1.01
Total	412	22 [5.3]	14	7	

*For one nymph, the specific strain wasn't identified

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