# Does getting defensive get you anywhere? - Seasonally varying selection in pea aphids shapes a dynamic infection polymorphism with a protective bacterial endosymbiont 

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#### Abstract

Facultative, heritable endosymbionts are found at intermediate prevalence within most insect species, playing frequent roles in their hosts' defense against environmental pressures. Focusing on Hamiltonella defensa, a common bacterial endosymbiont of aphids, we tested the hypothesis that such pressures impose seasonal balancing selection, shaping a widespread infection polymorphism. In our studied pea aphid (Acyrthosiphon pisum) population, Hamiltonella infection frequencies ranged from $23.2 \%$ to $68.1 \%$ across a six-month longitudinal survey. Rapid spikes and declines were consistent across fields, and we estimated that selection coefficients, for Hamiltonella-infected aphids, changed sign within this single season. Prior laboratory research suggested anti-parasitoid defense as the major Hamiltonella benefit, and costs under parasitoid absence. While a prior field study supported these forces as counter-weights in a regime of seasonal balancing selection, our present survey showed no significant relationship between parasitoid wasps and Hamiltonella. Field cage experiments provided some explanation: parasitoids drove ${ }^{\sim} 10 \%$ boosts to Hamiltonella frequencies that would be hard to detect under less controlled conditions. They also showed that Hamiltonella was not always costly under parasitoid exclusion, contradicting another long-held prediction. Instead, our longitudinal survey - and two overwintering studies - showed temperature to be the strongest predictor of Hamiltonella infection, matching some lab discoveries, and suggesting thermally sensitive costs and benefits, unrelated to parasitism, can shape this symbiont's prevalence. These results add to a growing body of evidence arguing for rapid, seasonal adaptation in multivoltine organisms. For many insects, such adaptation may be mediated through the diverse impacts of heritable symbionts on host phenotypes.


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C. Experimental design and results







