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Influence of delayed density and ultraviolet radiation on caterpillar granulovirus infection and mortality

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26

27 **Abstract**

- 28 1. Infectious disease is an important potential driver of population cycles, but this must occur
29 through delayed density-dependent infection and resulting fitness effects. Delayed density-
30 dependent infection by baculoviruses can be caused by environmental persistence of viral oc-
31 clusion bodies, which can be influenced by environmental factors. In particular, ultraviolet radi-
32 ation is potentially important in reducing the environmental persistence of viruses by inactivat-
33 ing viral occlusion bodies.
- 34 2. Delayed density-dependent viral infection has rarely been observed empirically at the popula-
35 tion level although theory predicts that it is necessary for these pathogens to drive population
36 cycles. Similarly, field studies have not examined the potential effects of ultraviolet radiation
37 on viral infection rates in natural animal populations. We tested if viral infection is delayed den-
38 sity-dependent with the potential to drive cyclic dynamics and if ultraviolet radiation influences
39 viral infection.
- 40 3. We censused 18 moth populations across nearly 9° of latitude over two years and quantified the
41 effects of direct and delayed density and ultraviolet radiation on granulovirus infection rate, in-
42 fection severity, and survival to adulthood. Caterpillars were collected from each population in
43 the field and reared in the laboratory.
- 44 4. We found that infection rate, infection severity, and survival to adulthood exhibited delayed
45 density-dependence. Ultraviolet radiation in the previous summer decreased infection severity,
46 and increased survival probability of the virus. Structural equation modelling found that the ef-
47 fect of lagged density on moth survival was mediated through infection rate and infection sever-
48 ity, and was 2.5 fold stronger than the effect of ultraviolet radiation on survival through infec-
49 tion severity.

50 5. Our findings provide clear evidence that delayed density dependence can arise through viral in-
51 fection rate and severity in insects, which supports the role of viral disease as a potential mech-
52 anism, among others, that may drive insect population cycles. Furthermore, our findings sup-
53 port predictions that ultraviolet radiation can modify viral disease dynamics in insect popula-
54 tions, most likely through attenuating viral persistence in the environment.

55 **Introduction**

56 Infectious disease plays an important role in population dynamics, potentially regulating
57 populations and driving cycles. Many insect populations undergo dramatic cyclic fluctuations, and
58 cyclic, delayed density-dependent disease outbreaks have been proposed as an explanation (Anderson
59 & May, 1980, 1981). However, many other mechanisms have also been proposed to explain insect
60 population cycles (Myers & Cory, 2013), particularly specialist parasitoids (Berryman, 1996). For
61 some species of Lepidoptera, empirical evidence has suggested that baculoviruses are most likely the
62 cause of cyclicality (Myers, 2000; Myers & Cory, 2013, 2016), though this has not been formally tested.

63 Delayed density-dependent feedbacks are required to drive cyclic population dynamics
64 (Turchin, 2003). Detection of delayed density-dependence involving many ecological factors has been
65 common (Turchin, 1990), and there have been several observations that viral epizootics follow high
66 densities of some insect species (Cory & Myers, 2003; Fuxa, 2004; Myers, 2000; Myers & Cory,
67 2016). However, direct demonstrations of delayed density-dependent viral infection rate and infection-
68 induced mortality are less common (Burthe et al., 2006; Fleming, Kalkmakoff, Archibald, & Stewart,
69 1986; Rothman, 1997). Observational studies over relatively small areas (>15km) using space-for-time
70 substitutions of local densities have shown delayed density-dependent incidence of viral infection in
71 voles (Burthe et al., 2006) and soil-dwelling hepialid caterpillars (Fleming et al., 1986). Experimental
72 manipulation of western tent caterpillar (*Malacosoma californicum pluviale*) densities at the tree level
73 showed delayed density-dependent infection rates by a nucleopolyhedrovirus (Rothman, 1997).

74 Besides host density, other aspects of the local environment may affect viral transmission and
75 dynamics in the field (Cory & Myers, 2003). In particular, ultraviolet radiation has been shown to
76 inactivate viruses of all kinds (Sagripanti & Lytle, 2007), including baculovirus occlusion bodies
77 (Griego, Martignoni, & Claycomb, 1985; Witt & Stairs, 1975). In field studies, examination of the
78 presence of baculovirus occlusion bodies on shaded vs. unshaded foliage suggested that sunlight on
79 leaves may inactivate viruses (Olofsson, 1988). A study of baculovirus transmission in forest tent
80 caterpillars (*Malacosoma disstria*) found depressed transmission rates on forest edges as opposed to the
81 forest interior, which was attributed to sunlight inactivating virus on leaves at forest edges (Roland &
82 Kaupp, 1995). A field study of two strains of gypsy moth NPV showed variable resistance of virus to
83 ultraviolet, in which inactivation rate from ambient ultraviolet was greater in a more potent strain than
84 a less potent one (Akhanaev et al., 2017). In another study of forest tent caterpillars using tree ring
85 analyses, it was suggested periods of weaker ultraviolet radiation increased the effect of forest tent
86 caterpillar outbreaks on tree growth (Haynes, Tardif, & Parry, 2018). In this example, less ultraviolet
87 radiation may have allowed better viral persistence and transmission and produced more severe
88 caterpillar outbreaks.

89 In the present study, we examined the effects of host density, delayed-density dependence, and
90 ultraviolet radiation on the survival, infection rate, and infection severity of an undescribed
91 granulovirus (see supplement) in Ranchman's tiger moth (*Arctia virginalis*). Ranchman's tiger moth
92 (Erebidae:Arctiinae) is a univoltine species with a long larval stage – usually lasting from late summer
93 until the following spring. Pupation occurs in late spring, and adults emerge in early summer, mate and
94 lay eggs during the summer. Larvae hatch mid-summer, and feed cryptically in litter and undergrowth
95 until they become larger in winter months. Caterpillars are generalists and feed on a variety of
96 herbaceous plants, with a preference for alkaloid-containing hosts (Karban, Karban, Huntzinger,
97 Pearse, & Cruisinger, 2010), and occur mostly in open grassland, shrubland, or savannah. Caterpillars
98 are parasitized by tachinid parasitoids, though caterpillars sometimes survive parasitism (English-Loeb,

99 Karban, & Brody, 1990). Parasitism has been found to have little role in population dynamics, leaving
100 the observed oscillatory dynamics largely unexplained (Karbon & de Valpine, 2010). Caterpillars also
101 sometimes show symptoms of granulovirus infection, which include poor growth, inactivity, watery,
102 opaque frass, regurgitation of milky fluid, and death. Granulovirus is a dsDNA virus that persists
103 outside of the host in the environment within a protective protein shell as an occlusion body (OB) .
104 Horizontal transmission can occur when caterpillars consume OBs on food sources, which dissolve in
105 caterpillars' alkaline gut and release infective virus particles (Vega & Kaya, 2012). Virus particles infect
106 the caterpillar starting from the epithelial tissue and then move to other parts of the body, including the
107 trachea, fat bodies, and hemolymph (Barrett, Brownwright, Primavera, & Palli, 1998), potentially
108 causing death of the caterpillar. In Ranchman's tiger moth (*Arctia virginalis*), granulovirus, like
109 parasitoids, does not uniformly kill its hosts.

110 To test the potential role of granulovirus in delayed-density dependent population dynamics in
111 Ranchmans' tiger moth and the influence of ultraviolet radiation on the persistence and transmission of
112 virus in moth populations, we censused 18 populations over two years along a gradient of latitude and
113 ultraviolet radiation and reared caterpillars from these populations in the laboratory. To identify the
114 mechanisms through which granulovirus affects moth population dynamics, we measured granulovirus
115 infection rate, severity, and survival of caterpillars to adulthood. We analyzed the effect of caterpillar
116 density and ultraviolet radiation at the study sites on viral infection and survival. We also tested if viral
117 infection mediated the effects of caterpillar density and ultraviolet radiation on caterpillar survival
118 using a structural equation model.

119 **Methods**

120 *Censuses*

121 Caterpillars were counted visually along 4 m wide linear transects of varying length (78-300m)
122 at 18 sites along a ~1000 km latitudinal gradient of the Pacific coast in California, Oregon, and
123 Washington (Figure 1). Transects were surveyed in 2019 and 2020, starting from 1 March through 30

124 May. We visited sites in each year from south to north, in accordance with the phenology of caterpillar
125 development so that caterpillars were surveyed when most were 4th or 5th instars (sampling dates in
126 Table S1). Transects were surveyed at a constant walking speed and all by the same observer (A. Pepi).
127 Density was estimated from caterpillar counts over the transect area (4m x length). During the second
128 year, sites were revisited within ten days from the calendar date of the first visit; however, this species
129 is a slow-growing caterpillar (roughly ten month development period), so density estimates were likely
130 not overly sensitive to small deviations in sampling date. In 2020, up to ca-. 30 caterpillars per site (or
131 less if fewer were found) were collected haphazardly and brought back to the laboratory for rearing.

132 *Rearing*

133 Caterpillars were reared individually in 6 oz plastic souffle cups with fabric lids and kept in an
134 incubator at 18 °C and 75% RH. Caterpillars were fed every 3-4 days with washed organic romaine
135 lettuce and yellow bush lupine (*Lupinus arboreus*) leaves collected from a part of Bodega Marine
136 Reserve without *Arctia virginalis*. Parasitoids that emerged from caterpillars were counted and
137 identified to family. Caterpillars that died during rearing were frozen at -20 °C for subsequent
138 dissection, except for those that clearly died due to emerged parasitoids. Caterpillars were reared until
139 pupation (~50 days), and pupae were placed together by site of origin (no more than ~10 per site) in 30
140 cm x 30 cm flight cages and allowed to emerge as moths, at room temperature. Pupae and moths were
141 reared until one month after the last moth emerged and sprayed once or twice weekly with water to
142 prevent desiccation. Adults do not feed. Dead pupae and moths were stored dry at room temperature
143 for subsequent dissection. Individuals that reached adulthood fully formed were counted as having
144 survived; individuals that died as caterpillars or failed to emerge from pupae or expand wings were
145 counted as not surviving.

146 *Dissection and viral assays*

147 After death, each individual was dissected to assess infection status. Two to six tissue samples
148 per insect were taken from the abdomen of adults and pupae or fat bodies of caterpillars, broken up

149 with forceps, smeared on a glass slide with water, and examined under a light microscope at 200x
150 magnification with phase contrast. Visible occlusion bodies (OB) ($<0.15\mu\text{m}$) were confirmed by
151 adding a drop of 1 M NaOH to the slide, which dissolves the viral protein coat, turning the OB
152 transparent (Lacey & Solter, 2012)(Figure S1). Tissue samples were taken until a positive NaOH test
153 was obtained; if no virus was found after six tissue samples, individuals were classified as uninfected
154 (see supplemental methods for more detail).

155 Infection severity was rated as uninfected, low, medium, high, or very high, based on the
156 number of samples required to discover virus, the density of OBs in tissue smear samples, and the
157 coloration of hemolymph and fat body (Figure S2). Low severity infections took many samples to
158 identify and had low densities of OBs. Medium severity infections took fewer samples to identify and
159 had moderate densities of OBs. High severity samples had infections in all samples with high densities
160 of OBs, with organ systems in the body still intact and identifiable. Very high severity infections had
161 infections in all samples with very high densities of OBs, with internal organ systems unidentifiable.

162 *Ultraviolet irradiation*

163 Daily values for ultraviolet radiation in 2019, specifically 'Erythemal Daily Dose' in Watts/m²
164 from OMI/Aura satellite data were accessed from NASA GES DISC (Hovila, Arola, & Tamminen,
165 2014). Values were extracted for each sampling location and the average value calculated from 360 to
166 180 days before the collection date of caterpillars (~March-September at most sites; see Table S1 for
167 collection dates). This timing coincided with the season during which viral OBs would have persisted
168 in the environment outside of hosts.

169 *Statistical analysis*

170 First, we analyzed the density-dependence of infection rate (proportion of infected individuals)
171 and the influence of ultraviolet radiation on infection rate. The proportion of all individuals (N=208)
172 that were infected was analyzed in response to log caterpillar density per m² from transects in 2019
173 (density in the previous year) and 2020 (density in the same year) using beta-binomial generalized

174 linear mixed models with site as a random effect. To test for the effect of ultraviolet radiation, we
175 implemented a model with both ultraviolet radiation and log density, to control for the effect of density
176 on infection rate.

177 Second, we analyzed the density-dependence of infection severity, and the influence of
178 ultraviolet radiation on infection severity. We analyzed infection severity (N=198 with severity rated)
179 in response to log density (same year and previous year) using cumulative link mixed models with site
180 as a random effect. As before, to test for the effect of ultraviolet radiation, we implemented a model
181 with both ultraviolet radiation and log density, to control for the effect of density on infection severity.
182 To test for effects of density and ultraviolet on infection severity independent of infection status, we
183 implemented models using only infected individuals (N=184). This allowed us to test whether different
184 processes influenced infection rate vs. infection severity. We implemented cumulative link mixed
185 models with only log density, and both log density and ultraviolet radiation as predictors as before.

186 Third, we tested for density-dependent survival and effects of ultraviolet radiation on survival.
187 We analyzed survival to emergence as fully formed moths (N=208) in response to caterpillar density,
188 tachinid parasitoid load, and ultraviolet radiation in a beta-binomial generalized linear mixed model
189 with site as a random effect.

190 Lastly, we tested our structural hypothesis that viral infection and infection severity were the
191 mechanisms through which lagged density and ultraviolet radiation affected survival in moth
192 populations. There are multiple possible pathways through which density, ultraviolet radiation,
193 infection rate, and infection severity might influence survival and thus population dynamics. To test
194 our specific structural hypothesis, we constructed a piecewise structural equation model which tested
195 whether infection rate and infection severity mediated the effects of ultraviolet radiation and caterpillar
196 density on survival. To do this, we transformed infection severity into a continuous variable between 0
197 and 1, by converting to numerical categories from 1-5, dividing by 5, and subtracting 0.1. We
198 constructed 3 component sub-models of our structural equation model: a beta-binomial model of

199 infection rate predicted by log density and ultraviolet radiation, a beta model of infection severity rate
200 predicted by infection rate, log density and ultraviolet radiation, and a beta-binomial model of survival
201 predicted by infection rate and infection severity (Figure 5). All sub-models included site as a random
202 effect, and only individuals that were not parasitized and were scored for infection severity were
203 included in the analysis (N=198). Standardized coefficients were calculated using the latent-theoretic
204 method (Grace, Johnson, Lefcheck, & Byrnes, 2018).

205 Beta-binomial and beta generalized linear mixed models were fitted using glmmTMB (Brooks
206 et al., 2017)□. Cumulative link mixed models were fitted using the ordinal package (Christensen,
207 2019). Piecewise structural equation models were fit using piecewiseSEM v. 1.2.1 (Lefcheck, 2016),
208 and component models were fit using glmmTMB. Ultraviolet radiation was logged and scaled to
209 improve model convergence. We checked for multicollinearity in models using VIFs;
210 multicollinearity was low to moderate in all models (<6) except for same year density in models of
211 infection severity (VIF from 10.5-11.5). Since we included same year density as a control variable, we
212 kept these models as is despite high VIFs. Plots were generated using ggplot2 (Wickham, 2009),
213 ggeffects (Lüdtke, 2018), and viridis (Garnier, 2018). All analysis and plotting were conducted in R
214 v. 3.6.3 (R Development Core Team 2020).

215 **Results**

216 *Infection rate*

217 The probability of infection increased with log density in the previous year in models of
218 infection rate (Table 1, Figure 2a). Infection rate had little relation to log density in the current year or
219 to ultraviolet radiation over the season when environmental OBs could have been exposed (Table 1,
220 Figure 2b).

221 *Infection severity*

222 Log density in the previous year increased infection severity in models of infection severity that
223 included both infected and uninfected caterpillars but this effect was weakened when ultraviolet

224 radiation was included in the model (Table 1, Figure 3a,b). This effect of ultraviolet radiation may
225 have been due to low power or to a negative correlation between log density in the previous year and
226 ultraviolet radiation (Pearson's r : -0.53). Ultraviolet radiation decreased infection severity when
227 included (Table 1, Figure 3c,d). In models that included only infected caterpillars, density in the
228 current and previous years had little effect on infection severity, and ultraviolet radiation had a negative
229 effect on infection severity (Table 1).

230 *Survival to adult emergence*

231 Survival to adult emergence decreased with log density in the previous year and with the
232 number of emerged parasites. Survival increased in response to density in the current year, and in
233 response to ultraviolet radiation (Table 1, Figure 4).

234 *Structural equation model*

235 We hypothesized a causal model linking ultraviolet radiation and disease to caterpillar survival
236 (Fig. 5). Shipley's d-separation test (Shipley, 2000) indicated that our structural equation model was
237 correctly specified (Fig. 5, Fisher's $C = 7.38$, $df = 6$, $P = 0.288$), meaning that there were no missing
238 paths in our causal model. The model results showed that the negative effect of log density in the
239 previous year on survival was mediated by infection rate and infection severity and that the effect of
240 ultraviolet radiation on caterpillar survival was mediated by infection severity (Fig 5). The relative
241 indirect positive effect size of log density in the previous year was ~2.5x greater than the positive effect
242 of ultraviolet radiation on survival (-0.085 vs. 0.035, calculated by multiplying path coefficients).

243 **Discussion**

244 Overall, our results show that log density during the previous year increased infection rate and
245 infection severity and decreased survival to adult emergence, suggesting a strong delayed density-
246 dependent effect of disease on caterpillar population dynamics. The effect of delayed density on both
247 infection rate and infection severity in univariate models and the result of the structural equation model
248 suggest that the effect of density on survival acts through infection severity as well as infection rate.

249 This is likely due to dose-dependent infection severity, in which caterpillars that are exposed to more
250 viral occlusion bodies develop more severe infections that result in death (Cabodevilla et al., 2011;
251 Eberle, Asser-Kaiser, Sayed, Nguyen, & Jehle, 2008; Matthews, Smith, & Edwards, 2002).

252 In contrast, log density in the current year, representing direct density-dependence, had no effect
253 on infection rate, a marginal negative effect on infection severity, and a positive effect on survival to
254 adult emergence. Though inconsistent, this positive effect may have been due to ascertainment bias:
255 sites that had larger viral outbreaks may have already declined in density over the course of the larval
256 season (summer 2019-spring 2020). Therefore, density at the time of collection had potentially a
257 reversed relationship to environmental OB density: sites that had declined from earlier high densities
258 would have higher levels of OB, resulting in more severe infections and a higher probability of death.
259 Overall, our results indicated negative delayed density dependence and weak or positive direct density-
260 dependence consistent with the second-order, oscillatory population dynamics observed in the system
261 over the long term (Pepi, Holyoak & Karban, unpublished analyses).

262 Ultraviolet radiation by contrast had no detectable effect on infection rate but reduced infection
263 severity and increased survival. The reduction in infection severity was even greater when only infected
264 individuals were considered. The effect of ultraviolet radiation on infection severity and survival but
265 not infection results may represent dose-dependent effects of exposure to OBs. Specifically, at sites
266 with higher ultraviolet radiation, more viral OBs were likely inactivated before caterpillars could be
267 exposed to them, so caterpillars received smaller doses of virus and thus became less severely infected
268 and were more likely to survive.

269 The observational nature of our study gives it some strengths and weaknesses. Our study is one
270 of few to show delayed-density dependent viral infection in the field, and the only study to show effects
271 of radiation on viral infection in a natural population of insects. Our results provide validation in a
272 field system to laboratory studies of virus dynamics and inactivation effects of ultraviolet radiation on
273 viruses (e.g., Bjørnstad et al., 1998; Witt & Stairs, 1975). Despite highly variable conditions between

274 sites, which were spread across a gradient of over 1000 km, we were able to detect clear effects of
275 delayed density on infection rate, severity, and survival to adulthood, and of ultraviolet radiation on
276 infection severity and survival to adulthood. We were also able to use structural equation modelling to
277 show that infection rate and severity mediate the effect of delayed density and ultraviolet radiation on
278 survival and strengthen the inferences made from our observational study.

279 Density in the previous year was a strong predictor of infection rates in populations (Fig 2).
280 This finding makes sense because there must have been a large enough population of hosts in the
281 previous year for the disease to spread and produce sufficient inoculum to persist into the current year.
282 Density in the previous year also affected infection severity (Fig 3), but this effect became much
283 weaker when ultraviolet radiation was included in models and even weaker when only infected
284 caterpillars were considered. In the structural equation model, infection severity was well explained by
285 infection rate because individuals must be infected to have high infection severity. This result may be
286 due to the conflation of infection rate and infection severity in the model since the same dose-
287 dependent mechanism may have caused them. However, in all models of infection severity that
288 included both ultraviolet radiation and delayed density, the effect of delayed density, beyond whether
289 individuals were infected or not, was weaker than the effects of ultraviolet radiation. Thus when effects
290 on infection severity were detected they may have in fact been generated by the spurious negative
291 relationship between delayed density and ultraviolet radiation. In contrast to infection severity,
292 ultraviolet radiation had no significant effect on infection rate. Ultimately, both delayed density and
293 ultraviolet radiation had strong and opposing effects on survival to adulthood, though the effect of
294 density in the previous year was much greater in magnitude. The effects of delayed density on survival
295 were mediated through infection with limited evidence of direct effect on infection severity. In contrast
296 the effects of ultraviolet radiation on survival were mediated entirely through effects on infection
297 severity.

298 A possible explanation for the finding that infection rates are primarily determined by host
299 density in the previous year but not by ultraviolet radiation is that viral infection may be determined in
300 part through vertical transmission (Burden, Griffiths, Cory, Smith, & Sait, 2002; Cabodevilla et al.,
301 2011; Cory, 2015; Cory & Myers, 2003). In particular, vertical transmission is likely to occur in *A.*
302 *virginalis* through persistent sub-lethal infections. We regularly observed OB in egg samples in adult
303 dissections, and thus eggs of adults with sublethal infections are likely contaminated with virus.
304 Vertical transmission represents a mechanism through which density in the previous year might
305 influence infection rates in the following year but is not subject to influence from ultraviolet radiation.
306 In this way, vertical transmission may maintain higher infection rates after high-density years with
307 epizootics, even if ultraviolet radiation attenuates inoculum in the environment (Cabodevilla et al.,
308 2011). Sublethal infections may represent another means by which granulovirus affects population
309 dynamics by imposing fitness costs on adults (Cabodevilla et al., 2011; Matthews et al., 2002;
310 Rothman, 1997), although we did not measure this in the present study.

311 In summary, we demonstrated population-level delayed density-dependent effects on viral
312 infection rate, infection severity, and survival to the reproductive stage, showing how viral infection
313 may drive cyclic dynamics in insects. We also showed for the first time that ultraviolet radiation may
314 influence disease dynamics and ultimately population dynamics of insects, through decreased infection
315 severity and increased survival rates. This provides support for the proposal that viral epizootics may
316 be an important mechanism driving cyclic dynamics of insects and Lepidoptera in particular (Myers &
317 Cory, 2013). Our findings also suggest that ultraviolet radiation may be an important factor to consider
318 as a driver of insect viral disease in the context of global change. In particular, long-term changes in
319 atmospheric transmittance of solar radiation, or “global dimming and brightening,” have been observed
320 and are potentially anthropogenic in origin (Wild, 2016). Such long-term changes may have the
321 potential to influence insect viral disease and population dynamics through changing levels of
322 ultraviolet attenuation of viral inoculum (Haynes et al., 2018).

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334 **Author contributions**

335 AP, VP, and RK conceived the study. AP and VP collected the data and conducted the analyses.
336 AP wrote the manuscript. All authors contributed critically to the drafts and approved final publication.

337 **Data Availability**

338 Data will be archived on KNB upon acceptance.

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473 **Figures & Tables**

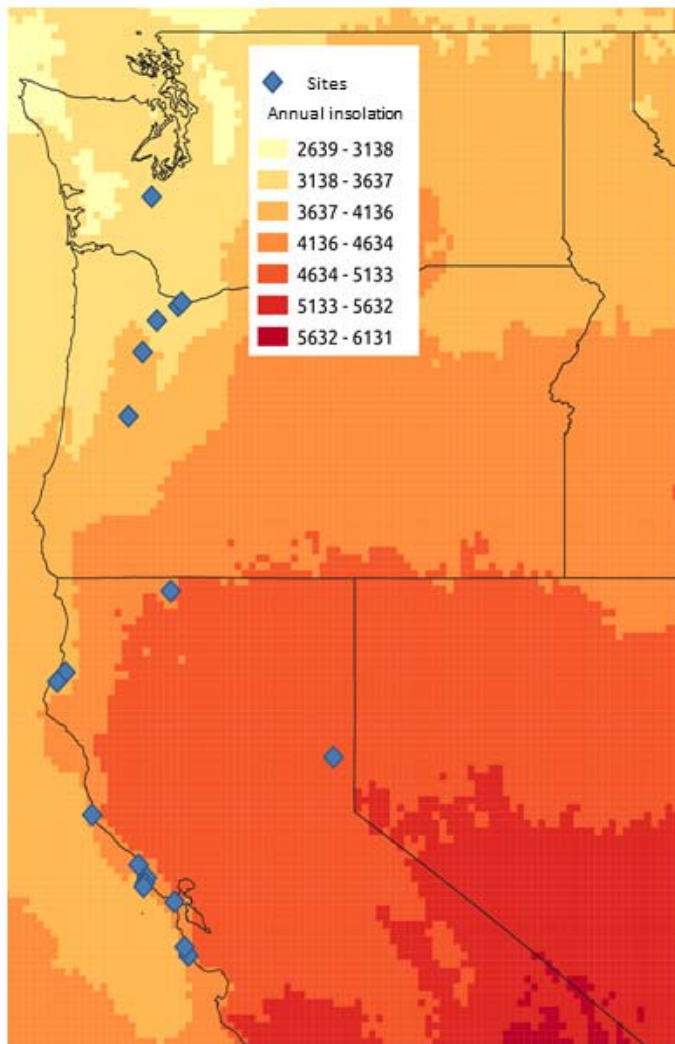
474 **Table 1.** Results of univariate models.

Model structure	Parameter	Logit $\beta \pm 1$ SE	z	P
Infection ~ log density	Log density current year	-0.20±0.38	-0.52	0.603
	Log density previous year	0.55±0.26	2.10	0.036
Infection ~ log density + ultraviolet	Log density current year	-0.10±0.39	-0.25	0.802
	Log density previous year	0.92±0.46	1.998	0.046
	Ultraviolet radiation	0.66±0.58	1.13	0.259
Severity ~ log density	Log density current year	-0.28±0.19	-1.49	0.135
	Log density previous year	0.26±0.11	2.35	0.019
Severity ~ log density + ultraviolet	Log density current year	-0.28±0.18	-1.61	0.101
	Log density previous year	0.16±0.11	1.37	0.170
	Ultraviolet radiation	-0.35±0.17	-2.04	0.041
Severity infected ~ log density	Log density current year	-0.27±0.19	-1.39	0.163
	Log density previous year	0.018±0.11	1.66	0.096
Severity infected ~ log density + ultraviolet	Log density current year	-0.28±0.18	-1.52	0.129
	Log density previous year	0.06±0.12	0.48	0.631
	Ultraviolet radiation	-0.44±0.18	-2.40	0.017
Survival ~ log density + ultraviolet	Log density current year	1.15±0.51	2.25	0.024
	Log density previous year	-0.75±0.31	-2.39	0.017
	Ultraviolet radiation	0.64±0.32	1.97	0.049
	Emerged parasites	-5.39±1.58	-3.42	0.0006

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479 **Figure 1.** Map of study site locations in Washington, Oregon, and California, overlaid on a map of
480 average annual insolation (GHI), in kWh/m².

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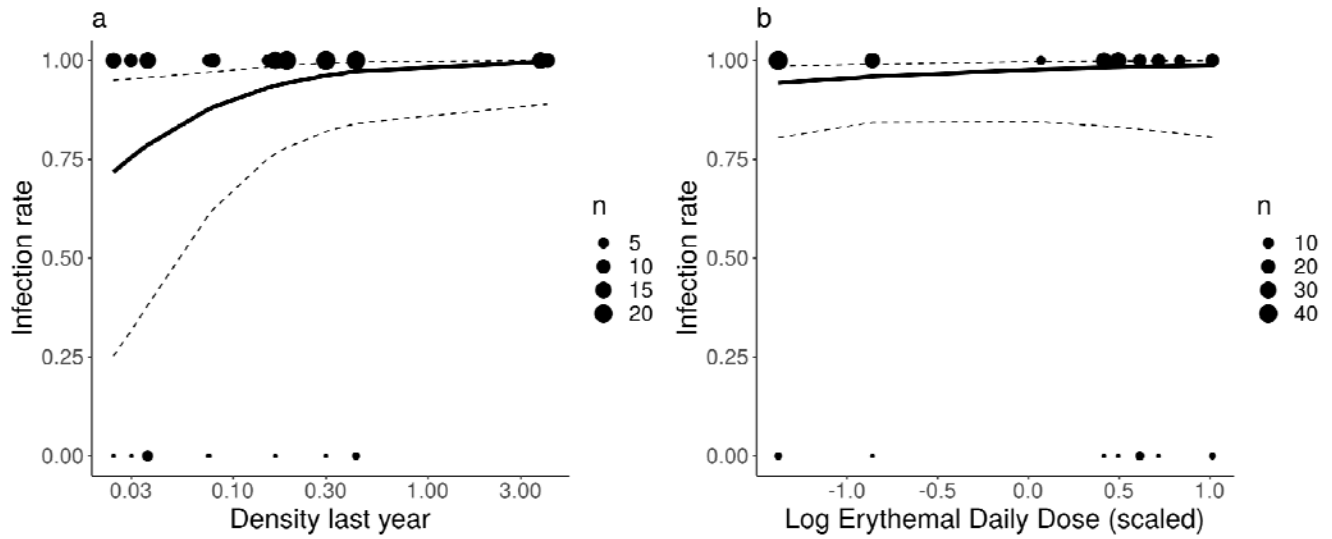
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489 **Figure 2.** Infection rate in response to (a) density in the previous year and (b) ultraviolet radiation as
490 scaled log erythemal daily dose in the summer before collection (2019) from beta-binomial generalized
491 linear mixed models.

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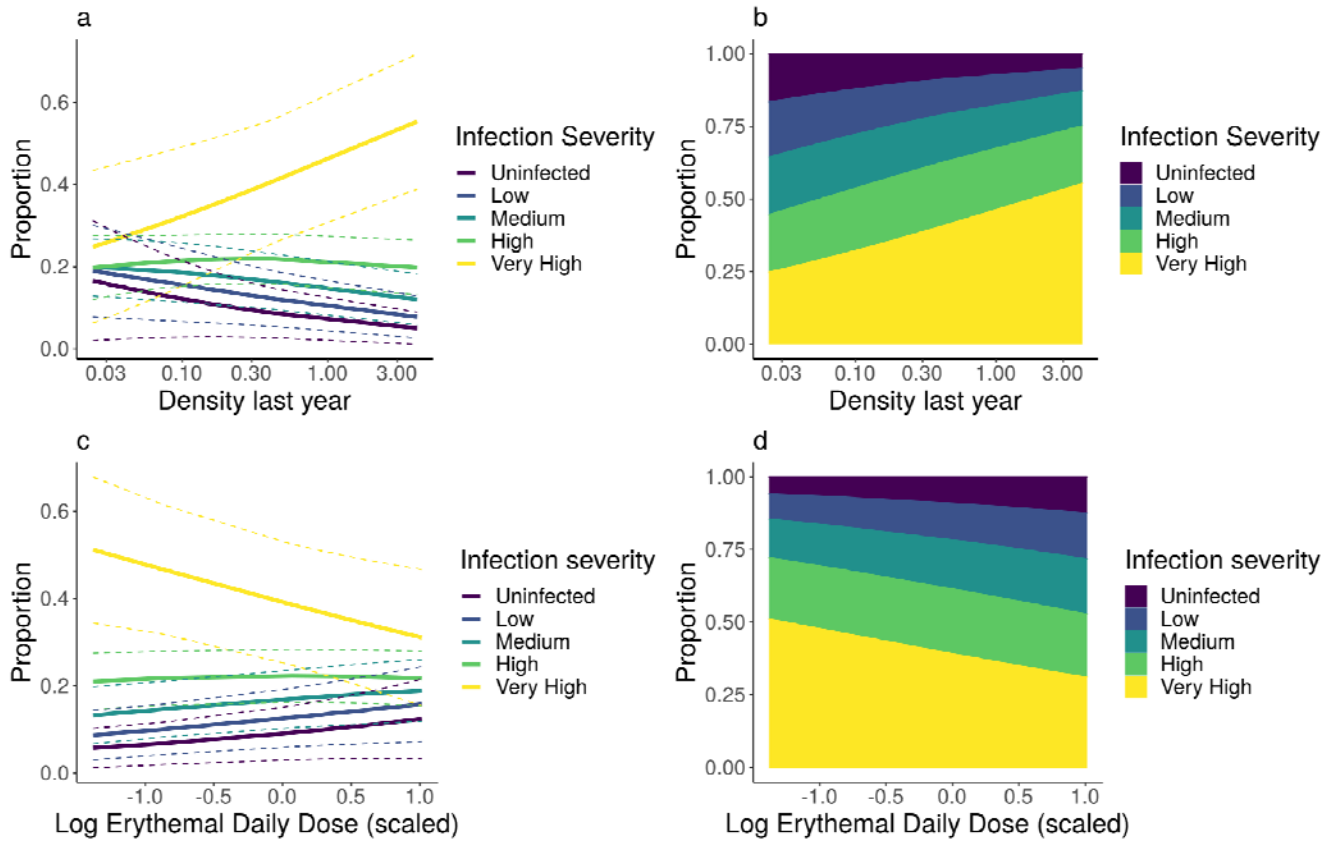
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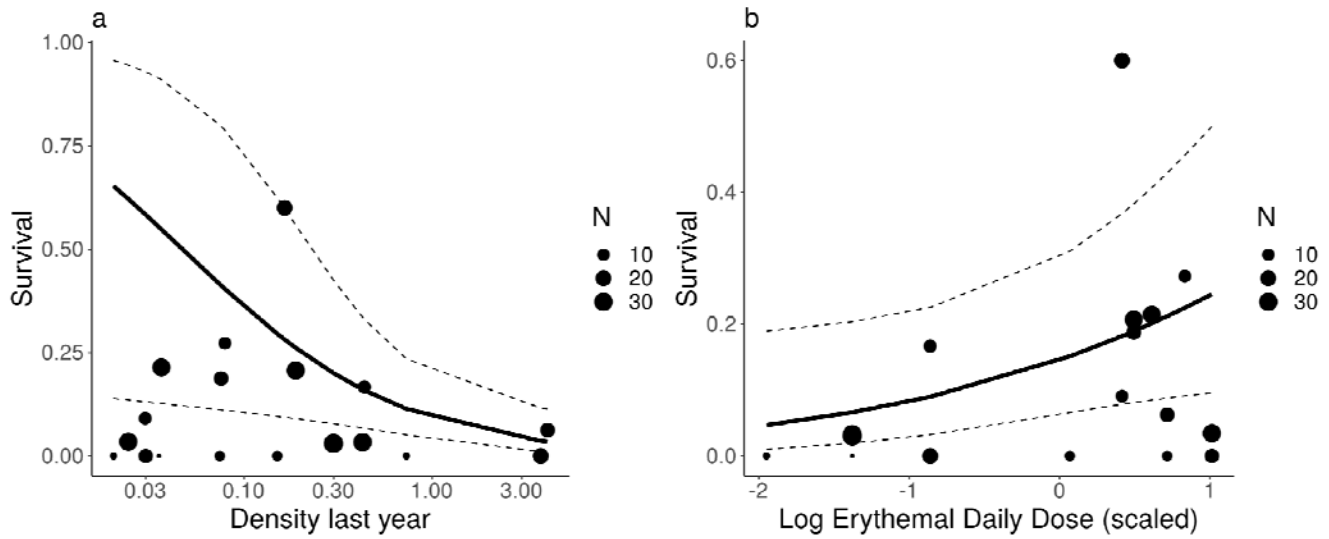
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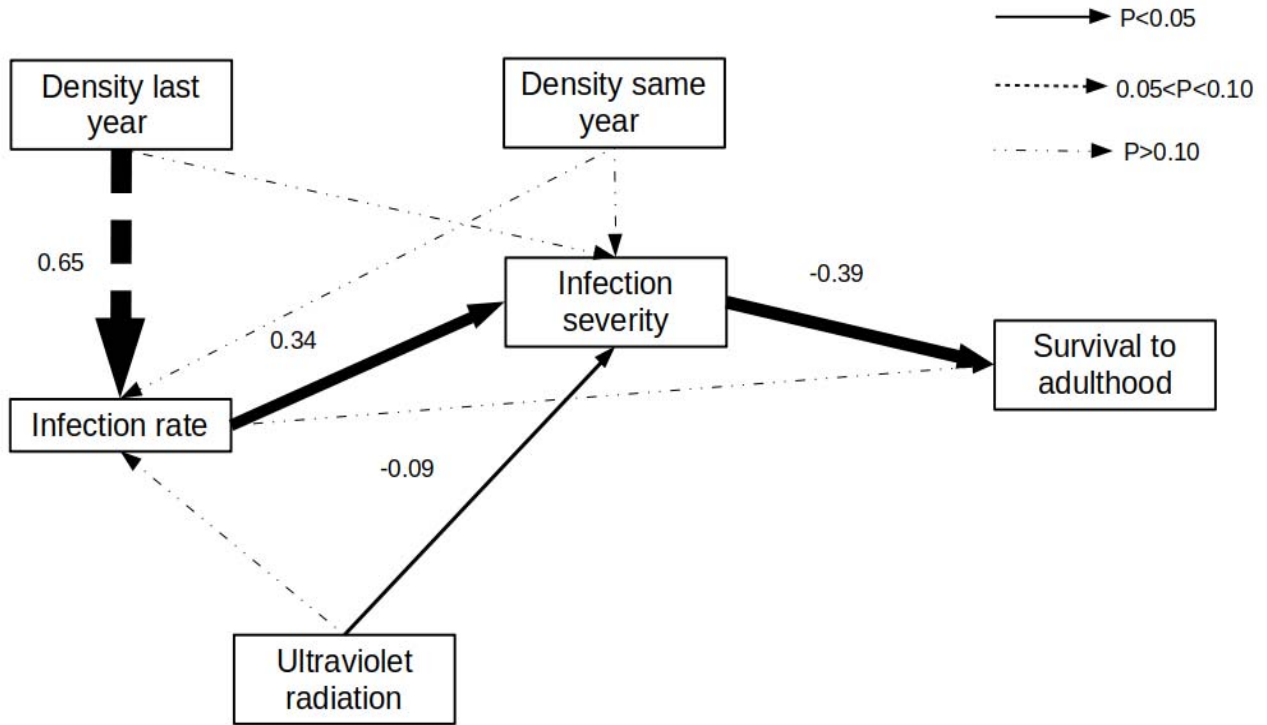
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Figure 3. Infection severity in response to (a-b) density in the previous year, and (c-d) ultraviolet radiation as scaled log erythemal daily dose in the summer before collection cumulative link mixed models. The left column (a,c) show model predicted rate of infection at each severity class with 95% confidence intervals, and the right column (b,d) shows the proportion of the total population predicted to be infected at each severity class by color.





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529 **Figure 5.** Path diagram of piecewise structural equation model. Size of arrows is proportional to the
530 standardized path coefficient (shown next to lines), and line type indicates P-value of the path
531 parameter.

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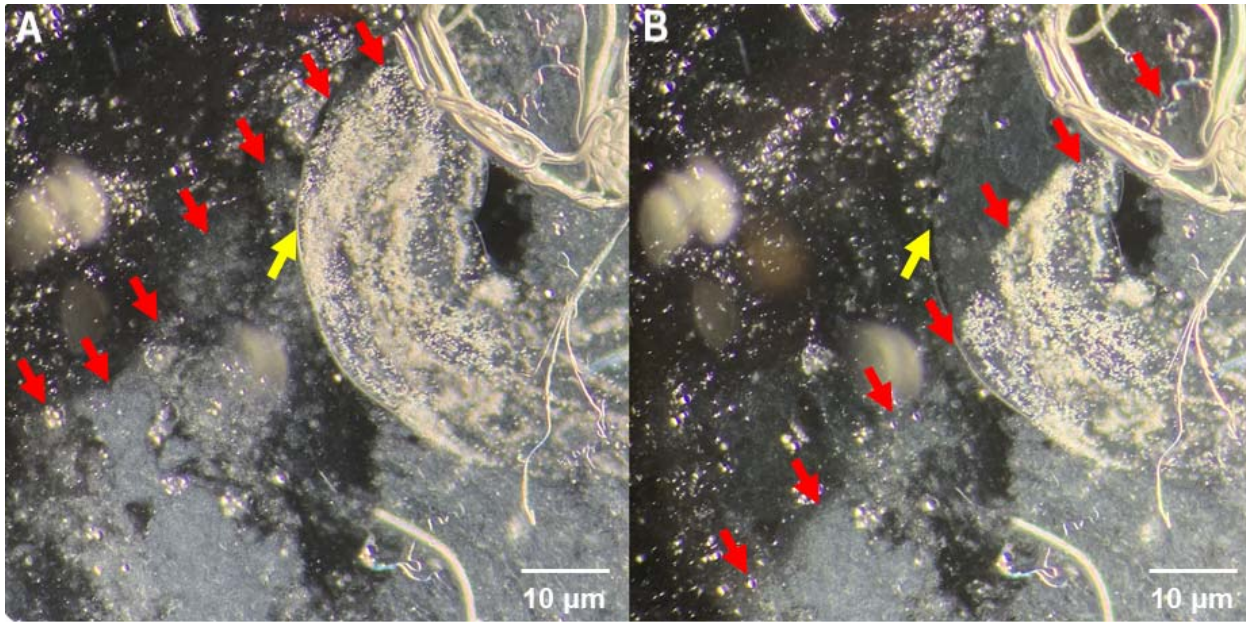
542 **Supplement**

543 **Table S1.**

Site	Latitude	Longitude	Number reared	Date Collected
Franklin Point, CA	37.155	-122.35	29	3/1/2020
Pescadero Marsh, CA	37.265	-122.41	16	3/1/2020
Golden Gate NRA, CA	37.844	-122.55	28	3/13/2020
Point Reyes South, CA	38.038	-122.99	15	3/13/2020
Point Reyes North, CA	38.088	-122.97	29	3/13/2020
Point Reyes Kehoe, CA	38.153	-122.94	2	3/13/2020
Bodega Marine Reserve, CA	38.319	-123.06	16	3/27/2020
Manchester State Park, CA	38.959	-123.72	6	3/27/2020
Loyalton, CA	39.700	-120.30	6	5/30/2020
Humboldt NWR, CA	40.670	-124.21	20	4/10/2020
Freshwater Farms, CA	40.786	-124.09	11	4/10/2020
Klamath River, Yreka, CA	41.829	-122.61	11	4/10/2020
Checkermallow Access, OR	44.072	-123.20	12	4/11/2020
Fairview Wetland, OR	44.895	-123.00	20	4/11/2020
Graham Oaks, OR	45.299	-122.80	30	4/11/2020
Powell Butte, OR	45.488	-122.50	33	4/11/2020
Salish Ponds, OR	45.531	-122.45	1	4/11/2020
West Rocky Prairie, WA	46.887	-122.87	2	4/12/2020

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547 **Figure S1.** Fat body tissue smear from a caterpillar infected with GV under a light microscope at 200x
548 magnification with phase contrast. A drop of NaOH was added which moved through the sample. The
549 red arrows point to the boundary between NaOH solution and water. The yellow arrows point to the
550 OBs (Figure A) which turned transparent (Figure B) after they were dissolved by the strong base.

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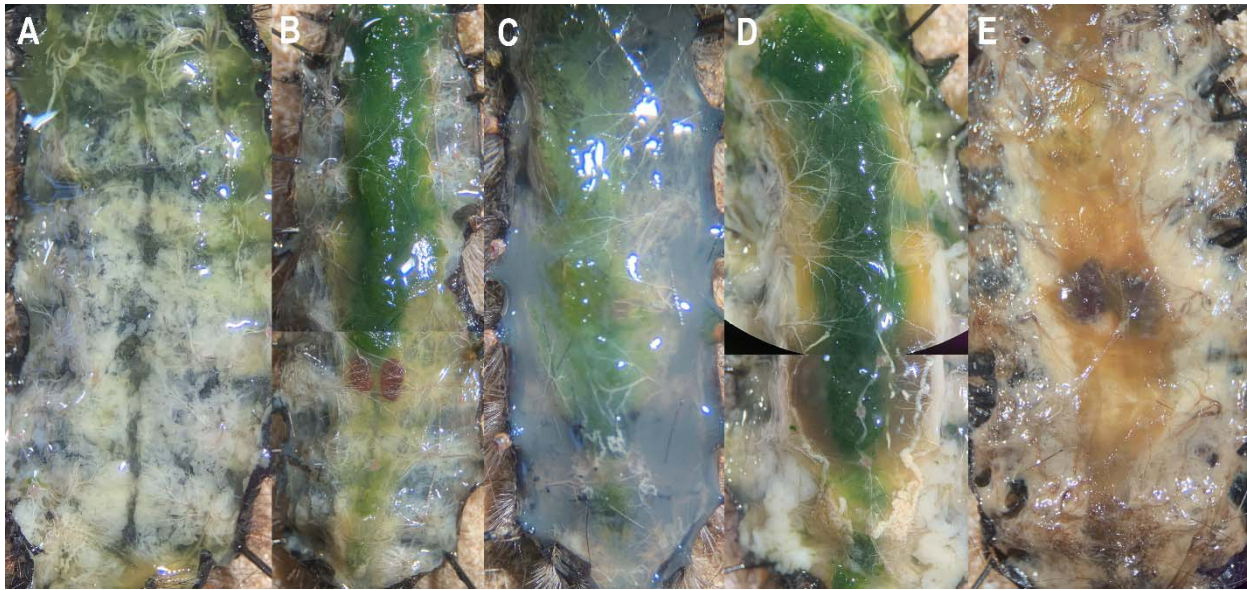
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562 **Figure S2.** Ventral dissection of late instar *A. virginalis* larva in the order of infection severity, ranging
563 from (A) uninfected, (B) low severity, (C) medium severity, (D) high severity, to (E) very high severity.

564 The caterpillar's gut in figure A is missing. Caterpillars with higher infection severity have a cloudier
565 hemolymph and a darker and more yellowish fat body. They also often have large clusters of infected
566 tissues next to the midgut (e.g. Figure D).

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