

Spruce budworm defoliation patterns during outbreak rise are influenced by tree species, insecticide spraying, and spatial autocorrelation

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7 insecticide spraying, and spatial autocorrelation

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19 **Abstract**

20 Spruce budworm (SBW; *Choristoneura fumiferana* Clem.) outbreaks are an important natural
21 disturbance in North America, killing trees over millions of hectares. We related 11 years of
22 SBW defoliation in 87 plots in Gaspé Peninsula, Québec, to 23 stand, site, and climate variables.
23 Defoliation was consistently ordered among host species: balsam fir > white spruce > black
24 spruce. Within the relatively small 200 km² study area, cluster analyses resulted in four and 10
25 clusters for balsam fir cumulative and current defoliation, respectively; variation in cumulative
26 defoliation converged over 11 years. Current defoliation was significantly spatially
27 autocorrelated among plots within stands, but autocorrelation weakened at distances >2500m.
28 Cumulative defoliation was significantly related to insecticide spraying, minimum and maximum
29 summer temperature, and interactions between SBW larvae per branch versus hardwood and
30 white spruce basal area. Tree species, insecticide spraying, and number of defoliating SBW
31 larvae were the main determinants of defoliation. Results showed much higher local spatial
32 variability in current defoliation patterns than previous studies, but over the course of an
33 outbreak, cumulative defoliation patterns converged. Cumulative defoliation patterns similar to
34 these, assigned based on local defoliation severity, can be input into defoliation-based growth
35 models to predict impacts on growth and survival.

36 **Keywords:** current and cumulative defoliation, mixed-effects modelling, cluster analysis, *Abies*
37 *balsamea*, *Picea glauca*

38 Introduction

39 Large-scale insect outbreaks are a major disturbance that impacts forests, reducing carbon
40 sinks, timber supply, and ecological services (Erdle and MacLean 1999). Outbreaks of spruce
41 budworm (*Choristoneura fumiferana* Clem., hereafter SBW) are the dominant natural
42 disturbance of balsam fir (*Abies balsamea* (L.) Mill.) and spruce (*Picea spp.*) forests in eastern
43 North America (e.g., Morris 1963; MacLean 2016). During an outbreak in the 1970s–1980s,
44 SBW caused defoliation covered ca. 50 million ha, with resulting losses of spruce-fir timber of
45 32–43 million m³ year⁻¹ (Sterner and Davidson 1982). Understanding how defoliation varies
46 across the landscape, as well as the underlying factors that shape these patterns, is essential for
47 developing strategies to manage forests in ways that reduce impacts associated with outbreaks
48 (MacLean et al. 2002; Hennigar et al. 2013).

49 Spatiotemporal patterns of SBW defoliation over large scales using aerial survey data have
50 been described in several studies. In Ontario, from 1941–1996 three distinct outbreak zones were
51 separated by two longitudinal corridors of lower defoliation severity (Candau et al. 1998). In
52 Québec from 1965–1992, cluster analysis revealed 25 unique defoliation patterns that explained
53 ~80% of the variation (Gray et al. 2000). For all of Canada from 1941–1998, Gray and
54 MacKinnon (2006) described 27 defoliation patterns, which they grouped into seven severity
55 classes ranging from “very mild” (<5 years of defoliation, average cumulative defoliation <50%)
56 to “severe” and “very severe” (>25 years of defoliation, average cumulative defoliation >300%).
57 Most recently, defoliation in Québec from 1938–2001 was reanalyzed by Cooke (2024), who
58 showed that the number of derived clusters may be arbitrarily high, depending on the selection
59 criteria used. Using 40 clusters, for example, identified behaviour that was “unique”: cycling in
60 the Ottawa valley at 24 years/cycle, as opposed to the 35-year cycle observed everywhere else
61 (Cooke 2024). Finally, in in New Brunswick from 1965–1992, Zhao et al. (2014) defined 28

62 defoliation patterns, grouped into five SBW outbreak classes. In all of these studies, defoliation
63 patterns differed in terms of duration, severity, and consistency, and were often found to be bi-
64 modal, with several years of low defoliation separating high-severity “peaks” (e.g., Gray and
65 MacKinnon 2006).

66 Several factors have been identified that significantly affect defoliation patterns. Because
67 of SBW’s relative specialization and feeding preferences, the availability of host species has a
68 particularly strong impact. Balsam fir is typically more defoliated than spruce, with white spruce
69 (*Picea glauca* (Moench) Voss), red spruce (*Picea rubens* Sarg.), and black spruce (*Picea*
70 *mariana* [Mill.] B.S.P.) in mixed fir-spruce stands sustaining 72%, 41%, and 28% as much
71 defoliation as balsam fir (Hennigar et al. 2008). This is consistent with the higher mortality rates
72 observed in balsam fir than spruce species (e.g., MacLean 1980; Nealis and Régnière 2004).
73 Forest hardwood content diminishes defoliation intensity, as stands with a higher proportion of
74 non-host hardwoods may foster a greater abundance and diversity of natural enemies (i.e.,
75 parasitoids and predators) (Su et al. 1996; Zhang et al. 2018) or increase larval dispersal losses
76 (Zhang et al. 2020) that reduces SBW populations. Insecticide spraying, if present in an area, is
77 used to reduce defoliation in the treatment year (Fuentelba et al. 2015, 2023); it is prevalent in
78 parts of eastern Canada and is important to account for in projecting effects of defoliation on
79 forest growth and survival. Elevation may influence defoliation (Bouchard and Auger 2014),
80 perhaps due to temperature differences or species composition variation with elevation.
81 Temperature affects tree vigor and therefore its ability to tolerate defoliation, and higher
82 mortality has been observed above 700m (e.g., Blais 1980; Witter 1985). Climate influences
83 defoliation patterns (Candau and Fleming 2005, 2011; Gray 2013), and affects autocorrelation of
84 defoliation spread (Bouchard and Auger 2014; Nenzén et al. 2018). Other factors, such as soil
85 drainage, slope, and aspect appear to be either less important or inconsistent in terms of their

86 impact on defoliation patterns (MacLean and Ostaff 1989; Bergeron et al. 1995).

87 Although species composition, insecticide spraying, elevation, and climate all influence
88 defoliation patterns, it can be difficult to distinguish their relative impacts from one another.
89 Climate, stand, and site data, for example, are often spatially autocorrelated i.e., locations close
90 to one another exhibit more similar values than those farther apart (Dormann et al. 2007). If
91 autocorrelation is overlooked, the assumption of data independence of residuals may be violated,
92 leading to false conclusions about the relative importance of different factors influencing
93 ecological relationships. Spatial autocorrelation analyses showed that that 28–47% of plots had
94 significantly clustered defoliation of trees within plots and a model including plot average
95 defoliation and subject tree basal area explained 80% of the variance (Li et al. 2019). A
96 regression tree was able to account for the spatial autocorrelation in defoliation frequency at all
97 but the shortest (<30 km) distances, where stand age or species composition and SBW dispersal
98 interact to overcome relationships with climate and forest characteristics (Candau and Fleming
99 2005). Fleming et al. (1999) observed that spatial autocorrelation waxed and waned through time
100 as the SBW went through sawtooth-like subcycles, probably as a function of variable density-
101 dependent dispersal. In survey data from New Brunswick, populations of second-instar larvae
102 were spatially aggregated throughout the declining phase of a SBW outbreak cycle but became
103 less so as densities decreased (Fleming et al. 1999). Ultimately, SBW defoliation is a function of
104 number of SBW larvae and population processes, most notably dispersal that is most likely to
105 explain time-varying spatial autocorrelation, but these may also be influenced by stand, site, and
106 climate.

107 Here, we analyse spatiotemporal defoliation patterns occurring over an 11-year period on
108 Gaspé Peninsula, Québec, beginning from 2012 when the outbreak was just starting to build up
109 in the area. Our data were derived from a network of 99 sample plots in the central area of the

110 Gaspé Peninsula, with annual defoliation assessments from 2011–2021 using more than 9,500
111 branch samples and 28,000 ocular tree defoliation assessments (Donovan and MacLean 2024).
112 These analyses extend past studies in several important ways. Whereas past studies tended to
113 focus on defoliation patterns occurring during the peak to declining phases of outbreaks, our
114 study may be the first to examine patterns during the early build-up phase of outbreak. Whereas
115 past studies relied mainly on relatively crude aerial survey data to construct patterns, our
116 analyses are based on precise branch-level data; aerial survey defoliation estimates in these plots
117 corresponded with mean branch sample defoliation in only 43% of cases and ranged from only
118 14%–58% in individual years (Donovan and MacLean 2024). Past studies were all at a
119 provincial or national scale, whereas our study area is only about 200km², but using precise
120 defoliation data to determine smaller scale defoliation patterns. Moreover, because there was an
121 ongoing foliage protection program in the area, we were also able to assess how periodic
122 insecticide treatments influenced defoliation patterns, which reflects reality in most eastern
123 Canada provinces.

124 Our study had three objectives. First, we used cluster analyses to classify plot-level
125 patterns of annual balsam fir, white spruce, and black spruce cumulative defoliation from 2011–
126 2021. Second, we assessed the occurrence of spatial autocorrelation in current-year balsam fir
127 defoliation patterns among plots within and between stands. Finally, we quantified the relative
128 importance of different factors, such as stand species composition, insecticide treatment history,
129 site, and climate variables on balsam fir current and cumulative defoliation.

130

131 **Methods**

132 **Study area and sample plot establishment**

133 The study area was located in the Gaspé Peninsula, Québec, part of the Temiscouata-

134 Restigouche Forest Section of the Great Lakes-St. Lawrence Region (Rowe 1972) (Fig. 1). SBW
135 defoliation was first detected in this region by annual aerial surveys in 2012 (MRNF 2021).
136 Forests in the area are composed mainly of balsam fir and white birch (*Betula papyrifera* Marsh.)
137 (Rowe 1972), with balsam fir and white spruce abundant at low elevations and black spruce
138 common in poorly drained areas. The landscape has rolling topography with rivers creating
139 narrow valleys and alluvial flats (Rowe 1972). Mean elevation is 365 metres (m), mean annual
140 temperature ranges from 0–2.5°C, mean annual precipitation from 900–1100 millimetres, and the
141 growing season lasts 150–160 days (Robitaille and Saucier 1998).

142 In 2014, 40 and 59 circular 0.04 ha sample plots were established in 28 stands northeast
143 and southwest of the towns of Amqui and Causapschal, respectively (Fig.1). All plots contained
144 balsam fir, white spruce, or black spruce (Supplementary Table S1); all Causapschal plots and
145 most Amqui plots were dominated by balsam fir, but 6 Amqui plots were selected to represent
146 hardwood-dominated or mixed hardwood-balsam fir (Zhang et al. 2018). Elevation of plots
147 averaged 284 m (range 177–404 m). Due to timber harvesting over the next decade, 12 plots
148 were lost, leaving 87 plots, averaging 165m apart, within 25 stands; there was an average of 3.5
149 plots per stand (range 1–6). Stands within the Amqui and Causapschal sites were within 4.3 km of
150 each other; and there was 40 km separating the Amqui and Causapschal locations (Fig. 1).
151 Therefore, we are examining defoliation at three scales: within stand, between stands, and
152 between the Amqui and Causapschal locations.

153 **Annual SBW defoliation measurements**

154 Percent current-year defoliation was measured in each plot in August after larval feeding
155 had finished by assessing shoots from one mid-crown branch sampled using pole pruners from
156 each of a target of 15 trees per host species per plot per year (Donovan and MacLean 2024).
157 Average height of dominant and co-dominant trees of the leading species in the stand (i.e.,

158 Lorey's mean height) ranged from 14.3m to 19.3m (Supplementary Table S1) and we usually
159 used 8–10 pole pruner sections (about 14–17 m) to sample branches, sometimes 11 sections (19
160 m) when required for the tallest trees. If 15 trees per host species were not present in a plot, we
161 sampled branches from all trees present; this occurred in 12% of balsam fir samples (where a
162 mean of 11 trees/plot were sampled), 43% of black spruce (mean 10 trees/plot sampled), and
163 66% of white spruce (mean 8 trees/plot sampled). The target sample sizes were based on
164 estimating mean stand defoliation with 90% confidence and $\pm 10\%$ error (MacLean and
165 MacKinnon 1998). Defoliation of current-, 1-, and 2-year-old foliage was measured on each of
166 25 randomly-selected shoots per branch, rated using seven percent defoliation classes (0, 1–20,
167 21–40, 41–60, 61–80, 81–99, 100%) (Fettes 1950). In 2014, the first year of sampling,
168 defoliation was estimated for older 2011, 2012, and 2013 foliage age-classes. Mean defoliation
169 per branch and foliage age-class was calculated using defoliation class mid-points. Mean
170 defoliation per host species per plot was calculated by weighting defoliation per tree by the basal
171 area of each sampled tree (Donovan and MacLean 2024). Mean cumulative defoliation per
172 species per plot over the 11-year period was calculated by summing current defoliation of that
173 species and plot for all years.

174 **Stand, site, and climate variables to be related to defoliation**

175 During plot establishment and again in 2018 and 2021, every tree ≥ 3 cm diameter at breast
176 height (DBH) was measured for species, DBH, height, height of base of live crown, and distance
177 and azimuth (degrees) from the plot centre. Heights and distances were measured with a Vertex
178 III and Transponder T3, Haglöf, Sweden. A total of 23 stand, site, or climate explanatory
179 variables was assembled for each plot (Table 1), based generally on the literature described in the
180 Introduction of factors thought to determine SBW defoliation levels, and were related to current-
181 year and cumulative defoliation. Seven stand explanatory variables included species composition

182 (basal area of balsam fir, white spruce, black spruce, hardwoods, and total basal area); annual
183 aerial spraying of the biological insecticide *Bacillus thuringiensis* spp. *kurstaki* (Btk) against
184 SBW (which began in 2015 and occurred in 11% of the samples, which henceforth we will term
185 plot-years), determined using spatially referenced spray blocks provided by the Québec
186 government. To be clear, the 87 plots were mostly sampled for 11 years each, but a few plots
187 were harvested and sampled for only 6–8 years; this resulted in a total sample of 819 annual plot
188 samples – i.e., 819 plot-years. Insecticide spraying occurred in a total of 87 plot-years, with 21
189 plots receiving at least 1 year of spraying: 6 plots received 1 or 2 years of spraying, and 15 plots
190 received 5 or 6 years of spraying. SBW previous-year overwintering second-instar larvae (L2)
191 populations were interpolated from the Québec government L2 sampling plot network, using the
192 inverse distance weighting interpolation tool in ArcGIS Pro version 3.2.2 (ESRI, 2023) with a
193 search neighbourhood of the L2 nearest estimates. Six site explanatory variables included
194 elevation, aspect, and slope measured directly in each plot; soil nutrient/moisture quality class
195 (poor, moderate, or rich soil nutrients and dry, moderate, or wet soil moisture), assigned based on
196 a soil pit dug and vegetation indicator species sampled in each plot and a site classification
197 system for the New Brunswick Kedgwick region (Zelazny et al. 1989), which is 25–32 and 60–
198 66 km from the Causapsal and Amqui plots, and part of the same Temiscouata-Restigouche
199 Forest Section (Rowe 1972) as our study area; and a depth-to-water index (Murphy et al. 2009)
200 calculated from a Québec LiDAR-derived digital elevation model (MRNF 2016) and the Arc
201 Hydro: Wetland Identification Model in ArcGIS Pro (Maidment 2002). Ten climate explanatory
202 variables included cumulative degree days $>5^{\circ}\text{C}$; summed monthly means per year of daily
203 extreme minimum and maximum temperatures for spring, summer, and winter (April-May, June-
204 July, November-March) and spring and summer precipitation (Table 1). These climatic periods
205 and variables were based on SBW biology and life stages, such as spring emergence larval

206 survival, summer larval feeding, and overwintering survival. The BioSIM (v11) Weather
207 Generator Editor (Régnière et al. 2017) was used to calculate 2011–2021 daily minimum and
208 maximum temperatures (°C) and precipitation (mm) for each plot based on latitude, longitude,
209 and elevation, using the four nearest weather stations, as recommended in the BioSIM manual.

210 **Cluster analysis of defoliation patterns**

211 Patterns of current and cumulative defoliation were grouped using the cluster analysis
212 unsupervised learning technique, which assesses values by virtue of their similarity to each other
213 and dissimilarity to other values (Everitt et al. 2011). Hierarchical cluster analysis used only
214 plots with all 11 years (2011–2021) of defoliation measurements: 54 plots for balsam fir, 9 for
215 white spruce, and 15 for black spruce. Yearly defoliation was standardized, centering means to
216 zero and scaled to one standard deviation, which ensures equal importance is given to any
217 defoliation year during clustering and is a common practice if variables have varying units and
218 ranges (James et al. 2021). The “scale” function in the R statistical language (R Core Team
219 2023) was used and graphing was done in RStudio version 2023.12.1.402 (R Core Team 2023).
220 We used the Euclidean distance matrix (James et al. 2021) computed using the “dist” function to
221 measure distances between pairs of values, the “hclust” function and “Ward.D2” linkage method
222 from the “stats” package (R Core Team 2023). The hclust function within R computes the actual
223 hierarchical clustering, and the Ward.D2 method (Ward 1963) is one option for controlling how
224 the algorithm creates clusters, minimizing within-cluster variance and helping to ensure data
225 within each cluster are similar to each other. To select the optimum number of clusters that best
226 described defoliation patterns, we examined three internal validation metrics: the total within-
227 cluster sum of squares (TWCSS), Dunn index (DI), and average silhouette width (SW). The
228 “clValid” and “factoextra” packages (Brock et al. 2008; Kassambara and Mundt 2020) were used
229 to calculate validation metrics for visual comparison for a range of 1 to 20 potential clusters. The

230 optimum number of clusters was heuristically selected when increasing the number of clusters
231 showed a negligible change based on averaging the three metrics.

232 **Spatial autocorrelation of defoliation patterns within and between stands**

233 Two spatial autocorrelation analyses were completed to characterize whether mean balsam
234 fir current defoliation of plots was clustered, dispersed, or random. The global Moran's I index
235 (Moran 1950) ranges from -1 to 1 and represents an overall measure of spatial autocorrelation
236 based on testing a null hypothesis that the attribute being examined has a complete spatial
237 randomness pattern among features for a study area. Accompanying p-values represent the
238 probability that an observed pattern is random, and z-scores represent the standard deviations
239 associated with a normal distribution. Global Moran's I analysis used the "Incremental Spatial
240 Autocorrelation" tool in ArcGIS Pro 3.2.2 (ESRI, 2023), testing a range of distances from 100–
241 600m in increments of 100m for within-stand, and 1000, 2500, 5000, 10,000, and 20,000m
242 distances to examine between-stand defoliation clustering.

243 The second spatial autocorrelation analysis determined the Getis-Ord G_i^* statistic, which
244 can help identify localized spatial clustering of neighbouring plots by identifying 'hot spot plots'
245 (highly defoliated plots surrounded by other highly defoliated plots) and 'cold spot plots' (lightly
246 defoliated plots surrounded by other lightly defoliated plots) (Getis and Ord 1992).

247 G_i^* equates to the sum of the differences between individual plots and the mean of all plots
248 representing a standard normal distribution z-score. When the local sum greatly differs from the
249 expected local sum based on random chance, statistically significant z-score results are obtained.

250 This analysis was conducted using the "Hot Spot Analysis" tool in ArcGIS Pro.

251 **Mixed-effects modelling of stand, site, and climate effects on balsam fir defoliation**

252 Differences in mean annual current defoliation among species were tested using Welch's
253 analysis of variance followed by pairwise comparisons using the Games-Howell method. This

254 method accounted for unequal variances and sample sizes among species and was conducted
255 using the “stats” and “rstatix” R packages (Kassambara 2023; R Core Team 2023).

256 We used generalized mixed-effects modelling (GLMM) for current defoliation (which is
257 bounded between 0–100%) and linear mixed-effects modelling (LMM) for cumulative
258 defoliation (which had a linear pattern year over year) to determine the effects of stand, site, and
259 climate explanatory variables (Table 1). Two- and three-way interaction terms were included to
260 test effects of significant and biologically relevant variable combinations on defoliation. GLMMs
261 defined with a binomial distribution and logit link function are suitable when response variables
262 are bounded percentage data (Zuur et al. 2009). Both GLMMs and LMMs enable correct
263 inference when handling complex data that are hierarchical in nature, have non-independence,
264 and include missing observations (Zuur et al. 2009). Models dealt with hierarchical and repeating
265 observations, reducing spatiotemporal autocorrelation by defining random intercepts for sample
266 plots nested within stands and crossing with year as the random effect structure. All suitable
267 fixed-effect explanatory variables were standardized to have a mean of zero and scaled to one
268 standard deviation, enabling easier interpretability of model estimates. Assumptions (e.g.,
269 residual homogeneity, linearity) were checked, and modelling used the “glmer” and “lmer”
270 functions in the “lme4” package (Bates et al. 2015).

271 Explanatory variables were initially tested separately and ranked in importance based on
272 parameter estimates, t-statistic (LMM) or z-statistic (GLMM), and p-value. Subsequent
273 modelling based on maximum likelihood estimations started with the most significant variable
274 and iteratively added (forward selection) the next most significant variable, resulting in the
275 largest improvement of model goodness-of-fit (James et al. 2021; Zuur et al. 2009). At each
276 model iteration adding an additional variable, analysis of variance likelihood ratio testing,
277 Akaike Information Criterion (AIC), Bayesian Information Criterion (BIC), marginal R^2 ,

278 conditional R^2 , and root mean squared error (RMSE) were evaluated; final models were
279 identified based on the lowest BIC. Pearson's pairwise collinearity matrix (Supplementary Fig.
280 S1) followed by iterative checks of variance inflation factors (VIFs) helped identify signs of
281 multicollinearity; VIFs >10 were considered an issue (Dormann et al. 2013), and analyses
282 ensured VIFs were <5 for selected variables. Final GLMM log-odds estimates were converted to
283 probabilities to better understand the effects of fixed-effect variables on the response.

284

285 **Results**

286 **Defoliation variability among species and plots**

287 Annual current and cumulative defoliation for all plots combined by host species are
288 presented as box plots in Fig. 2. Current defoliation increased each year from 2011 to 2015,
289 reached a maximum in 2015 and 2017, and then was lower in at least some plots from 2018 to
290 2021 (Fig. 2a). Although there was wide variability among plots, demonstrated by the “length”
291 of box plots in Fig. 2a, consistent trends among species were evident. “Sawtooth-like
292 fluctuations” (sensu Royama 1984) were evident, with local peaks for white spruce and balsam
293 fir in 2015, 2017, and 2020. The order of defoliation was always balsam fir $>$ white spruce $>$
294 black spruce (Fig. 2a). Mean defoliation of balsam fir was significantly greater than white spruce
295 in 7 years (2011–2014 and 2016–2018) (letters above the x-axis in Fig. 2a). In the other 4 years,
296 defoliation of white spruce was almost as high as balsam fir (Fig. 2a). In contrast, mean
297 defoliation of black spruce was significantly lower than balsam fir in all years and lower than
298 white spruce in 7 of 11 years (Fig. 2a). The highest mean current defoliation of black spruce in
299 all plots was about 30%, compared to about 50% for white spruce and 70% for balsam fir.
300 However, maximum current defoliation of individual plots was 100% for balsam fir, about 80%
301 for white spruce, and 60% for black spruce (Fig. 2a).

302 Mean cumulative defoliation, calculated as the sum of current annual defoliation, gradually
303 increased over the 11 years and reached nearly 500% for balsam fir, 350% for white spruce, and
304 150% for black spruce (Fig. 2b). These values can be interpreted as complete removal of the
305 equivalent of about 5, 3.5, and 1.5 age-classes of foliage for the three species, respectively.
306 Maximum cumulative defoliation among all plots over the 11 years was 650% for balsam fir,
307 400% for white spruce, and 200% for black spruce (Fig.2b).

308 **Cumulative defoliation patterns of balsam fir and spruce**

309 Cluster analysis divided the 54 plots with complete 2011–2021 balsam fir defoliation data
310 into four clusters, with 9 to 20 plots per cluster (Fig. 3a). Clustering validation metrics led us to
311 select four clusters, based mainly on SW but also TWCSS and DI metrics (Supplementary Fig.
312 S2). Clusters 1 to 4, respectively, had mean cumulative defoliation of balsam fir after 11 years of
313 362%, 411%, 549%, and 615% (Fig. 3a). Three years earlier, in 2018, the four clusters had the
314 same order of mean cumulative defoliation. Cluster 1 had the lowest cumulative defoliation and
315 the highest frequency of insecticide spraying (29 plot-years, in 7 of the 11 years) (Fig. 3a).
316 Cluster 2 also had a high frequency of insecticide spraying (24 plot-years, in 6 years). Cluster 3
317 was the most common pattern, and only had 7 plot-years of insecticide spraying. Cluster 4 had
318 no insecticide spraying. Temporal patterns of cumulative defoliation were generally similar
319 among clusters, except Clusters 1 and 2 had lower defoliation before 2015 and substantially
320 lower defoliation due to insecticide spraying (Fig. 3a). Clusters 2 and 3 were the dominant
321 defoliation patterns at the Causapscal site, whereas all four clusters were present at the Amqui
322 site (Fig. 3a).

323 Although balsam fir was the dominant species (Supplementary Table S1), there were
324 sufficient white spruce in five stands and black spruce in six stands to conduct cluster analyses
325 on spruce (Fig. 3b, 3c). Most previous regional studies have combined defoliation patterns for fir

326 and spruce species. White spruce had four clusters, culminating in about 220%, 310%, 325%,
327 and 400% mean cumulative defoliation, respectively (Fig. 3b). Defoliation in the three clusters
328 for black spruce resulted in mean defoliation levels of about 120%, 150%, and 220% (Fig. 3c).
329 Four of the seven spruce clusters resulted from plots in single stands. Four white spruce plots and
330 four black spruce plots received aerial applications of Btk in 2–7 years (Fig. 3b, 3c).

331 **Current annual defoliation patterns of balsam fir**

332 Cluster analysis results were more complex for current defoliation (Fig. 4). We selected 10
333 clusters as optimal, based on diminishing returns when averaging the TWCSS, DI, and SW
334 validation metrics for various numbers of clusters (Supplementary Fig. S2). In contrast to
335 cumulative defoliation, current defoliation clusters tended to represent plots in only one or two
336 stands, with only clusters 8 and 10 representing plots from 3 or 4 stands (Fig. 4). Insecticide
337 spraying against SBW occurred in only 5 of the 10 clusters and coincided with years of low
338 defoliation in some cases but not in others (Fig. 4). Although cumulative defoliation smoothly
339 increased over time (Fig. 3), year-to-year fluctuations determined the current defoliation cluster
340 results (Fig. 4). Plots in the most severe cumulative defoliation cluster 4 (Fig. 3a) were all in
341 current defoliation clusters 9 or 10 (Fig. 4). Clusters 1, 2, and 5–7 represented plots from the
342 Causapscal site, whereas clusters 3, 4, 8, and 9 had plots from both study sites, and cluster 10
343 contained only plots from the Amqui site (Fig. 4). Cluster 1 (stand 16) was late to the outbreak
344 and was located somewhat distal from the others (Fig. 1). Defoliation in most clusters declined in
345 2016, but not in Cluster 3, which was noteworthy in having high hardwood content in stand 25
346 and high black spruce in stand 10 (Supplementary Table S1).

347 With few exceptions, the 2–5 plots within each stand had very similar balsam fir current
348 defoliation patterns over time (Supplementary Fig. S3). Only two of the 24 stands (numbers 15
349 and 24) had plots in more than one current defoliation cluster (Fig. 4). This suggests that stand-

350 to-stand variability in current defoliation was much greater than plot-to-plot variability within
351 stands, and sampling effort would be more productive with more individual stands than
352 replicated plots within stands.

353 **Spatial autocorrelation of balsam fir current defoliation within and between stands**

354 Based on global Moran's I results, spatial clustering of balsam fir current defoliation within
355 a given stand was statistically significant in all years and distances (Table 2). Moran's I was
356 highest at 0.94 in 2016, the year in which defoliation decreased sharply, and lowest but still
357 significant in 2012 at 0.41 (Table 2). Across the within-stand distances tested (100–600m), 100–
358 300m often (71% of the time) had the highest Moran's I in any given year (Table 2), suggesting
359 mean defoliation severity generally was similar at short distances among plots. At between-stand
360 distances, plots were significantly clustered at 2500 and 5000m distances in all years, but
361 Moran's I values were lower than within stands (Table 2). Defoliation was not significantly
362 clustered for 2 of the 11 years at 5000m and for 5 years at distances of 10000 and 20000m (Table
363 2).

364 Both significant 'hot spot' (highly defoliated) and 'cold spot' (lightly defoliated) plots
365 occurred (Fig. 5). The proportion of hot spot plots ranged from 7–25% among years (Table 3),
366 generally with mean current defoliation >60%, whereas cold spot plots ranged from 6–25%, with
367 mean current defoliation <50% (Fig. 5). Although current defoliation was significantly clustered
368 in all years, it really occurred spatially at low or high defoliation levels, and 54–80% of plots
369 (range among years) were not significantly clustered (Table 3; Fig. 5). Hot and cold spot plots
370 did occur within the same stands in a given year, indicating that current defoliation was often
371 spatially autocorrelated (mapped examples are shown in Supplementary Fig. S4).

372 **Stand, site, and climate factors influencing SBW defoliation patterns**

373 Results of forward-selection modelling of explanatory variables affecting balsam fir

374 cumulative and current defoliation are shown in Table 4. The LMM that we selected for balsam
375 fir cumulative defoliation included five significant main effect variables: Sprayed, SuMinTp,
376 SuMaxTp, HWba, and WSba, and interactions between L2branch*HWba, and L2branch*WSba
377 (Model M4b; Table 4). Adding interaction terms of L2branch with HWba and WSba*L2branch
378 slightly lowered AIC and BIC (Table 4), resulting in significant yet minor effects on cumulative
379 defoliation (Fig. 6f and 6g). Fixed effects accounted for 9% of cumulative defoliation variation
380 (marginal R^2), and random effects explained an additional 84% variation (conditional R^2 ; 2% for
381 plots nested within stands, 7% for stands, and 83% for year variable) (Table 4).

382 Spraying insecticides resulted in a strong negative effect (Sprayed, -84.1; Table 5),
383 reducing cumulative defoliation by 34–177% each year from 2015–2021 (Fig. 6a). Summed
384 summer monthly minimum temperature had a weaker negative effect (SuMinTp, -19.3; Table 5);
385 early in the outbreak (2011–2016), cumulative defoliation increased as SuMinTp increased, but
386 later this shifted to a negative relationship (Fig 6b). Summed summer monthly maximum
387 temperature showed a positive but inconsistent effect (SuMaxTp, 29.8; Table 5) on cumulative
388 defoliation; positive in 4 years, negative in 2 years, and no relationship in 4 years (Fig. 6c). Main
389 effects of hardwood (HWba, -17.1) and white spruce (WSba, -10.4) basal area both negatively
390 affected defoliation (Table 5; Figs. 6d and 6e). The interaction of L2branch*HWba negatively (-
391 15.2; Table 5) affected cumulative defoliation; for moderate SBW levels, 4 of 11 years were
392 negative and 3 years positive, whereas 3 of 6 years having high SBW were negative (Fig. 6f).
393 The interaction of L2branch*WSba was also negative (-8.5; Table 5) but inconsistent; for
394 moderate SBW, 5 of 9 years were negative and 3 years positive, while for higher SBW levels, 4
395 of 6 years were negative (Fig. 6g).

396 The selected GLMM for balsam fir current defoliation included six significant main effect
397 variables: Sprayed, L2branch, WiMaxTp, SuMaxTp, SuMinTp, HWba, and interactions of

398 Sprayed*SuMinTp, and L2branch*HWba (Model M5b, Table 4). The interaction term for HWba
399 in most years indicated a stronger effect in reducing current defoliation at high L2 levels (Fig.
400 7g) and that current defoliation of sprayed plots decreased with SuMinTp but at low defoliation
401 levels (Fig. 7f). The current defoliation model had a marginal R^2 of 0.19 and conditional R^2 of
402 0.54 (2% for plots, 8% for stands, and 90% for grouping variable year) (Table 4).

403 Annual spraying of insecticides had a strong negative effect on current defoliation (log-
404 odds values are shown in Table 5), with mean current defoliation reduced by 11–60% due to
405 spraying in 6 years (Fig. 7a). 2017 was the exception, with mean current defoliation 9% greater
406 in sprayed than unsprayed plots (Fig. 7a). Overall, only 11% of the 819 plot-years analyzed were
407 sprayed with insecticide, with spraying concentrated in specific stands or clusters from 2015 to
408 2021 (Figs. 3, 4, and S3). L2branch positively affected current defoliation, once mean defoliation
409 exceeded 25% in 2013 (Fig. 7b). SuMinTp negatively but inconsistently affected current
410 defoliation, with defoliation increasing with warmer summer minimum temperatures in 5 years,
411 versus decreasing in 5 years (Fig. 7c). SuMaxTp also negatively but inconsistently affected
412 current defoliation, negative in 5 years and positive in 5 years (Fig. 7d). WiMaxTp positively
413 affected current defoliation in 7 of 11 years (Fig. 7e). The interaction of Sprayed*SuMinTp was
414 inconsistently negative but indicated spraying reduced current defoliation at lower severities for
415 3 of 6 years from 2015–2021 (Fig. 7f). Finally, the interaction of L2branch*HWba showed a
416 negative reduction of current defoliation. Defoliation decreased with increasing HWba in 4 of 9
417 years with moderate SBW and in 3 of 6 years with high SBW levels (Fig. 7g).

418

419 Discussion

420 Our study provides a unique examination of the spatiotemporal defoliation patterns that
421 occur during the buildup of a SBW outbreak, as well as identifying some of the key factors

422 driving these trends. Cluster analysis of balsam fir current-year defoliation simplified patterns
423 into 10 clusters, reflecting varying annual fluctuations among years within plots and stands. In
424 contrast, cluster analysis of cumulative defoliation converged into only four patterns. This has
425 implications for simplifying determination of impacts of SBW outbreaks, since cumulative
426 defoliation is strongly related to growth reduction and tree survival (e.g., Erdle and MacLean
427 1999; Hennigar and MacLean 2010). For example, local sampling of SBW defoliation could be
428 used to infer one of the four cumulative defoliation patterns we observed, or other higher or
429 lower cumulative defoliation, and used as input into existing defoliation-based stand growth
430 models (e.g., Erdle and MacLean 1999) to determine tree growth reduction, mortality, and
431 volume yield using models or a decision-support system (e.g., MacLean et al. 2001, 2002).
432 Results showed that defoliation patterns can vary considerably within our small geographic study
433 area, in contrast to the large regional defoliation patterns described in previous studies (e.g.,
434 Candau et al. 1998; Gray and MacKinnon 2006; Zhao et al. 2014) and that defoliation differed
435 among host species that were combined in previous studies.

436 Tree host species significantly affected current and cumulative defoliation severities, with
437 balsam fir > white spruce > black spruce, as found in previous studies (Hennigar et al. 2008;
438 Bognounou et al. 2017). Defoliation of white spruce was closer to that of balsam fir during
439 severe outbreak years, as also noted by Hennigar and MacLean (2010), Fuentealba et al. (2019),
440 and Bauce et al. (2024).

441 In any given year, within-stand spatial autocorrelation of balsam fir current defoliation was
442 significant when the severity of defoliation was light or severe but generally not when it was
443 moderate. This is probably a function of effects of SBW dispersal and high larval survival
444 driving population eruptions. When a major moth influx occurs, high SBW populations occur on
445 most trees or plots and result in similarly high (severe) defoliation in the area. When populations

446 are low, resulting in light defoliation, SBW only occurs close to oviposition sites on individual
447 trees, so plots in an area have similarly light defoliation.

448 Balsam fir is the primary host for SBW due to the phenology synchronization between
449 diapause emergence and bud flush during spring (Nealis and Régnière 2004), and possibly also
450 due to white spruce having superior defensive chemicals (e.g., Delvas et al. 2011). In the years
451 with highest balsam fir defoliation (2015, 2019–2021), white spruce had levels of defoliation
452 approaching that of balsam fir. Bognounou et al. (2017) referred to this as associational
453 susceptibility, where secondary hosts can temporarily become more susceptible from an influx of
454 SBW populations, optimal climate perturbations, or primary resource depletion leading to
455 spillover to secondary hosts. Our data also showed evidence of an associational resistance
456 interaction between black spruce and balsam fir, similar to Bognounou et al. (2017). Black
457 spruce was common throughout the Causapsca site and had low defoliation levels throughout
458 the years sampled, even when balsam fir in the same stands sustained severe defoliation.

459 Our cluster analysis of defoliation patterns was unique compared to previous studies
460 because it was based on precise branch defoliation data (Donovan and MacLean 2024), and the
461 spatial scale focused on patterns within and between stands for a small part of the SBW range. In
462 contrast, previous studies used less precise aerial survey data (Donovan and MacLean 2024)
463 covering provincial or national scales (Candau et al. 1998; Gray and MacKinnon 2006; Zhao et
464 al. 2014). Even at these different spatial scales, there were similarities in results. Clusters for
465 current annual defoliation consistently resulted in various distinctive “up-and-down, sawtooth”
466 patterns, whereas cumulative defoliation patterns were categorized by only a few groups (Gray
467 and MacKinnon 2006; Zhao et al. 2014). Such similarities across temporal (i.e., different
468 outbreaks) and spatial scales (i.e., different locations) provide support for use of cumulative
469 defoliation scenarios in analyses of impacts using a decision-support framework (e.g., MacLean

470 et al. 2002). Although our defoliation patterns were from a small $\sim 200\text{km}^2$ area, results showed
471 that 10 current annual defoliation patterns converged over time into only four cumulative
472 defoliation patterns. Local SBW defoliation patterns derived from aerial surveys or satellite
473 imagery can be used to assign these of other cumulative defoliation patterns to specific
474 stands/forest and used as input into defoliation-based stand growth models (e.g., Erdle and
475 MacLean 1999) to predict growth reduction, mortality, and guide harvest replanning, salvage,
476 and insecticide prioritization to minimize losses (e.g., MacLean et al. 2001, 2002; Hennigar and
477 MacLean 2010). Cluster analysis of defoliation patterns also showed the strong effect of aerial
478 spraying of insecticides in reducing defoliation, which occurred only in 11% of the plot-years,
479 confirming their effectiveness in forest protection (Fuentelba et al. 2015).

480 Balsam fir current defoliation patterns showed consistent synchrony within and between
481 nearby stands based on spatial autocorrelation analysis. Many studies of forest insects link spatial
482 autocorrelation at different scales to the Moran effect (Peltonen et al. 2002; Royama 2005).
483 Population synchrony is driven by favourable spatially autocorrelated climate conditions and
484 insect dispersal capabilities (Nenzén et al. 2018; Larroque et al. 2019). An essential aspect of
485 climate effects on SBW dynamics relates to temperature sensitivity of all life stages (Pureswaran
486 et al. 2015). Our analysis identified SuMinTp, SuMaxTp, and WiMaxTp as having positive and
487 negative effects on observed defoliation, indicating that climatic differences over multiple years
488 may play a significant role in defoliation differences.

489 Our hot and cold spot analysis indicated that within-stand spatial autocorrelation occurred
490 at both light and severe defoliation levels. This suggests that multiple fine-scale interactions at
491 the stand level likely support or counteract the broad-scale dispersal and climate mechanisms
492 leading to localized SBW population fluctuations. There was generally a high proportion of plots
493 showing significant clustering at both defoliation $>80\%$ and $<\sim 40\%$, but these thresholds varied

494 among years, especially for 2017. The hot and cold spot analysis essentially showed that
495 defoliation of plots within a stand or in nearby stands tend to be clustered when defoliation was
496 high and when it was low, but not intermediate. This might reflect effects of dispersal-driven
497 sawtooth defoliation patterns but also high larval survival resulting in high defoliation, or the
498 opposite for low defoliation level cold spots. In a previous study using a subset of our data, Li et
499 al. (2019) found that spatially clustered defoliation also occurred among trees within plots,
500 usually under high mean plot defoliation. Density-dependent regulation of SBW interacting with
501 natural enemies (e.g., parasitoids, diseases, and predators) (Royama 1984; Bouchard et al. 2018)
502 can contribute to both within-plot and local between-plot spatial clustering of defoliation.

503 Our mixed effect modelling strategy used forward selection methods and checking model
504 goodness-of-fit and variance explained. Some simpler models performed almost as good as our
505 final selected models. Careful consideration was given to additional main effects and interaction
506 terms and how increasing model complexity helped explain variability in defoliation. For
507 example, including the interaction of SBW L2branch*HWba significantly improved model
508 goodness-of-fit but resulted in a marginal reduction in overall error, and a slight decrease in
509 variance explained for current defoliation. In the context of SBW population dynamics, this
510 relationship is an important mechanism due to increased natural enemy abundance and diversity
511 with increased hardwood composition resulting in lower SBW populations and defoliation (Su et
512 al. 1996; Zhang et al. 2018). The interaction of SBW L2branch*WSba is also relevant since
513 white spruce offers adequate survival for SBW populations compared to balsam fir but is prone
514 to less defoliation due to spruce's increased ability to produce numerous shoots (Nealis and
515 Régnière 2004). However, from a management standpoint, insecticide spraying, L2 population
516 level, and species composition can be used to estimate defoliation patterns.

517 Although our models of current and cumulative defoliation were significant, there were
518 large differences between marginal R^2 , which explains the variance for only fixed effects, and
519 conditional R^2 , which explains model variance for both fixed and random effects. The “Year”
520 variable random effects explained a lot of the model variability and indicated that previous year
521 defoliation was important to subsequent year defoliation, due to temporal autocorrelation.

522 Aerial spraying of insecticides had the strongest effect of any factors in reducing balsam fir
523 current and cumulative defoliation. Spraying insecticides is an effective protective strategy for
524 keeping host trees alive and reducing defoliation severity (Fuentelba et al. 2015, 2023). This
525 effect was also evident in our cluster analysis of defoliation patterns as lower severity patterns
526 were associated with more frequent insecticide spraying. Number of SBW L2 had a positive
527 effect, causing increases in current defoliation, as would be expected. Number of SBW L2
528 (overwintering) are used as the threshold for triggering an early intervention insecticide use
529 strategy against SBW (Johns et al. 2019; MacLean et al. 2019), which aims to minimize
530 defoliation by detecting and treating localized rising SBW populations and also under a foliage
531 protection strategy (Fuentelba et al. 2023).

532 Even though it is a cold-adapted species (Pureswaran et al. 2015), SBW overwintering
533 survival is negatively influenced by extreme cold conditions, with a combination of temperature
534 and exposure durations required to kill half of the population (LTT_{50}) of 11.4 hours at -39°C
535 (Delisle et al. 2022). Although SBW larvae can withstand very low temperatures in winter, they
536 are vulnerable to cold spells during early diapause or post-diapause development (Delisle et al.
537 2022). Extreme winter warming can also deplete SBW larval energy reserves, resulting in
538 immediate or delayed mortality (Han and Bauce 1998; Moise et al. 2024). The threshold
539 temperature triggering post-diapause development is thought to be near $2.5\text{--}5.5^{\circ}\text{C}$ (Bean 1961;
540 Miller et al. 1971), and temperatures near this threshold may be one reason for higher mortality

541 during spring post-diapause development (Han and Bauce 1998). Overwintering survival is
542 further complicated by host species selected by SBW (Berthiaume et al. 2020; Moise et al.
543 2023). Winter temperature fluctuations and timing are critical, and acute warming periods during
544 early diapause reduce survival in contrast to warming periods later in diapause (Moise et al.
545 2024). Our results suggest that WiMaxTp was within the threshold range since there was a
546 positive effect on current defoliation patterns. Interestingly, during 2021, we observed an
547 unseasonably warmer winter than in previous years, which caused a negative effect on
548 defoliation patterns. Candau and Fleming (2005, 2011) also found that warmer winter minimum
549 and maximum temperatures were associated with higher defoliation frequency.

550 SuMinTp and SuMaxTp were significant factors, but their effects on defoliation patterns
551 were inconsistent between years. Warmer SuMinTp negatively affected current and cumulative
552 defoliation. An interaction between spraying and SuMinTp negatively affected current
553 defoliation, which might reflect less effective spraying or operational issues achieving optimal
554 timing under cold temperatures. However, it is important to recognize that we used coarse
555 seasonal climatic measures (e.g., seasonal annual minimum and maximum temperatures) over 11
556 years, so any relationships with insecticide efficacy or SBW population dynamics are
557 speculative. Warmer SuMaxTp negatively affected current defoliation and positively affected
558 cumulative defoliation. Several previous studies have found somewhat similar results. Gray
559 (2013) found that higher SuMinTp and SuMaxTp resulted in longer but less severe outbreaks,
560 with the mechanism hypothesized to be potential disruption to natural enemies' survival
561 dynamics along with reduced SBW feeding rates (Bellemin-Noël et al. 2021). Blais (1958) and
562 Régnière et al. (2012) observed that cooler summer temperatures contributed to slower SBW
563 larval development and reduced survival. Both Candau and Fleming (2011) and Nenzén et al.
564 (2018) conducted SBW outbreak projections, finding that summer temperatures along with

565 spring and winter temperatures were important. Increasing SuMaxTp up to a threshold of 18°C
566 resulted in the increased probability of severe defoliation for a SBW outbreak in Newfoundland
567 (Zhang et al. 2023). Such results among literature and our study suggest there are multiple
568 compounding factors related to temperature causing complexity of SBW feeding and survival
569 during the summer period.

570

571 **Conclusions**

572 Host tree species strongly influenced SBW current and cumulative defoliation over an 11-
573 year period, with balsam fir > white spruce > black spruce, similar to results from most previous
574 studies. On average, balsam fir defoliation was 13% greater than white spruce and 29% greater
575 than black spruce.

576 Cumulative defoliation tended to converge into a few general patterns, which cluster
577 analysis placed into 4, 4, and 3 patterns for balsam fir, white spruce, and black spruce,
578 respectively. In contrast, balsam fir current defoliation had much more year-to-year fluctuation,
579 which cluster analysis grouped into 10 patterns. The ability to aggregate cumulative defoliation
580 patterns can assist in building accurate geo-referenced defoliation-based growth reduction
581 estimates based on stand characteristics and stand-to-stand defoliation spread and
582 synchronization.

583 Spatial autocorrelation of balsam fir current defoliation within and between stands was
584 significant, especially at low and high defoliation severities, and gradually weakened when
585 distances exceeded >2500m. From an outbreak monitoring viewpoint, future ground-based
586 defoliation assessments would be more efficient by deploying 1–3 sample plots per stand, to
587 increase the number of assessed stands.

588 Stand species composition and insecticide treatment were the primary determinants of

589 defoliation patterns, with climate showing minor effects and site no effects. Significant factors
590 related to balsam fir current and cumulative defoliation patterns were Sprayed, L2branch,
591 SuMinTp, SuMaxTp, WiMaxTp, and interactions L2branch*HWba, and Sprayed*SuMinTp.
592 Spray and L2branch were expected variables included in final models, since spraying reduces
593 SBW populations and defoliation, and overwintering L2branch estimates are a proxy for the
594 following year feeding larval SBW population. SuMinTp had a negative effect on defoliation but
595 was inconsistent among years; SuMaxTp had negative and positive effects on current and
596 cumulative defoliation, respectively; and WiMaxTp had a positive effect on current and
597 cumulative defoliation patterns. Each interaction term had a negative effect on current and
598 cumulative defoliation, but significance was marginal and inconsistent across years. Such
599 relationships, which are known to influence SBW larval development, phenology, overwintering
600 mortality, and host-natural enemy interactions, suggest that temperature can also significantly
601 influence defoliation patterns over a long period within a relatively small local area of <200km².

602

603 **CRedit author contribution statement**

604 **Shawn D. Donovan:** Project conceptualization, Data curation, Formal analysis, Visualization
605 Writing – original draft, review & editing. **David A. MacLean:** Project conceptualization,
606 Supervision, Administration and funding acquisition, Writing – original draft, review & editing.
607 **Chris Hennigar:** Writing – review & editing. **Rob C. Johns:** Writing – review & editing. **Yun**
608 **Zhang:** Writing – review & editing.

609 **Declaration of Competing Interest**

610 The authors declare that they have no known competing financial interests or personal
611 relationships that could have appeared to influence the work reported in this paper.

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621 **Data Availability**

622 Data generated or analyzed during this study are available from the corresponding author upon
623 reasonable request.

624

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849 **Table 1.** Description of 23 plot-level explanatory variables (fixed effects) grouped by stand, site
 850 and climate categories examined for their influence on balsam fir current and cumulative
 851 defoliation patterns using mixed effects models. Model random effect structure included sample
 852 plots nested with stands and year as a crossed random effect.

	Explanatory Variables	Description	$\bar{\chi}$	σ
Stand	Sprayed	Annual aerial spraying of insecticides		
	L2branch	Spatially interpolated estimates of SBW L2 populations per branch	19.5	13.9
	BFba	Balsam fir basal area (m ² /ha)	26.9	12.2
	WSba	White spruce basal area (m ² /ha)	2.8	4.8
	BSba	Black spruce basal area (m ² /ha)	5.6	9.7
	HWba	Hardwood basal area (m ² /ha)	7.1	8.9
	Totalba	Total basal area (m ² /ha)	43.2	6.8
Site	Elevation	Ground elevation above sea-level (m)	284	65
	Aspect	Compass direction of downward slope (0–360 degrees)	239	90
	Slope	Inclination of downward slope (0–90 degrees)	5.2	3.8
	RootDep	Rooting depth (3 classes <20cm, 20–40cm, >40cm)		
	SoilQual	Soil nutrient/moisture quality (6 classes used out of 9 possible classes)		
	DTW	Depth-to-water index (0–55, lower values equate to water near surface)	11.9	11.7
Climate	SpCDD	Spring cumulative degree days >5°C from Apr.–May	133.7	32.5
	SuCDD	Summer cumulative degree days >5°C from Jun.–Jul.	663.0	41.5
	SpMinTp	Summed spring monthly minimum temperature from Apr.–May	-19.2	3.4
	SuMinTp	Summed summer monthly minimum temperature from Jun.–Jul.	6.4	2.8
	WiMinTp	Summed winter monthly minimum temperature from Nov.–Mar.	-124.9	10.0
	SpMaxTp ^a	Summed spring monthly maximum temperature from Apr.–May	44.2	4.8
	SuMaxTp ^b	Summed summer monthly maximum temperature from Jun.–Jul.	59.4	3.2
	WiMaxTp ^a	Summed winter monthly maximum temperature from Nov.–Mar.	34.1	9.3
	SpPrcp	Summed spring monthly precipitation (mm) from Apr.–May	182.1	43.7
	SuPrcp ^b	Summed summer monthly precipitation (mm) from Jun.–Jul.	188.3	52.1

853 * Sprayed, L2branch and all climate variables include annual plot-level observations.

854 ^{a,b} The same superscripts indicate pair-wise collinearity ($r > 0.70$) between explanatory variables.

855 **Table 2.** Overall spatial autocorrelation trends at varying distances for balsam fir current
 856 defoliation per plot within stands (100–600m) and between stands (1000–20,000m) based on
 857 global Moran's *I*. Bolded values indicate highest results per year and asterisks signify not
 858 rejecting null hypothesis due to randomness of defoliation patterns.

Balsam fir current defoliation global Moran's <i>I</i> by distance (m)											
Year	Plots within stands						Plots between stands				
	100	200	300	400	500	600	1000	2500	5000	10000	20000
2011	0.48	0.46	0.44	0.43	0.44	0.46	0.43	0.37	0.23	0.22	0.22
2012	0.41	0.57	0.58	0.55	0.54	0.52	0.45	0.45	0.30	0.25	0.25
2013	0.69	0.77	0.75	0.74	0.70	0.66	0.53	0.29	0.05	0.04	0.03
2014	0.82	0.81	0.80	0.80	0.76	0.72	0.63	0.21	0.01*	-0.02*	-0.02*
2015	0.83	0.85	0.85	0.85	0.84	0.85	0.84	0.51	0.08	-0.02*	-0.02*
2016	0.93	0.94	0.94	0.93	0.93	0.91	0.87	0.63	0.33	0.17	0.17
2017	0.83	0.74	0.71	0.69	0.64	0.58	0.50	0.42	0.30	0.20	0.20
2018	0.86	0.85	0.82	0.81	0.75	0.65	0.45	0.22	0.03*	0.01*	0.02*
2019	0.83	0.84	0.83	0.84	0.84	0.84	0.81	0.72	0.25	0.05	0.05
2020	0.70	0.84	0.85	0.82	0.81	0.80	0.78	0.63	0.20	-0.01*	-0.02*
2021	0.85	0.88	0.87	0.86	0.85	0.85	0.81	0.72	0.51	0.01*	-0.02*

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860 **Table 3.** Spatial autocorrelation Getis-Ord G_i^* analysis of balsam fir current defoliation per
 861 year: proportion of plots labelled as ‘hot spots’ (highly defoliated plots surrounded by other
 862 highly defoliated plots) and ‘cold spots’ (lightly defoliated plots surrounded by other lightly
 863 defoliated plots), or not significantly clustered ($\alpha = 0.05$) using a maximum within stand plot
 864 distance of 600m. Number of plots (n) per year declined over time due to timber harvesting.

Balsam fir current defoliation hot/cold spot analysis of sample plots (%)				
Year	n	Hot spots	Cold spots	Not clustered
2011	84	14	6	80
2012	84	20	10	70
2013	84	19	13	68
2014	84	25	21	54
2015	84	13	23	64
2016	84	14	25	61
2017	75	7	21	72
2018	75	17	12	71
2019	57	18	23	59
2020	54	6	17	77
2021	54	19	19	62

866 **Table 4.** Generalized and linear mixed models (GLMM, LMM) examining relationships between balsam fir current and cumulative
 867 defoliation (response variables) and site, stand, and climate explanatory variables. Model analyses using maximum likelihood estimation
 868 applied stepwise forward explanatory variable selection considering model goodness-of-fit Akaike and Bayesian Information Criterion
 869 (AIC, BIC), and analysis of variance likelihood ratio tests. Marginal and conditional R² and root mean squared error are based on
 870 restricted maximum likelihood estimation. **Bolded** and *italicized* text identifies the final models selected based on lowest BIC.

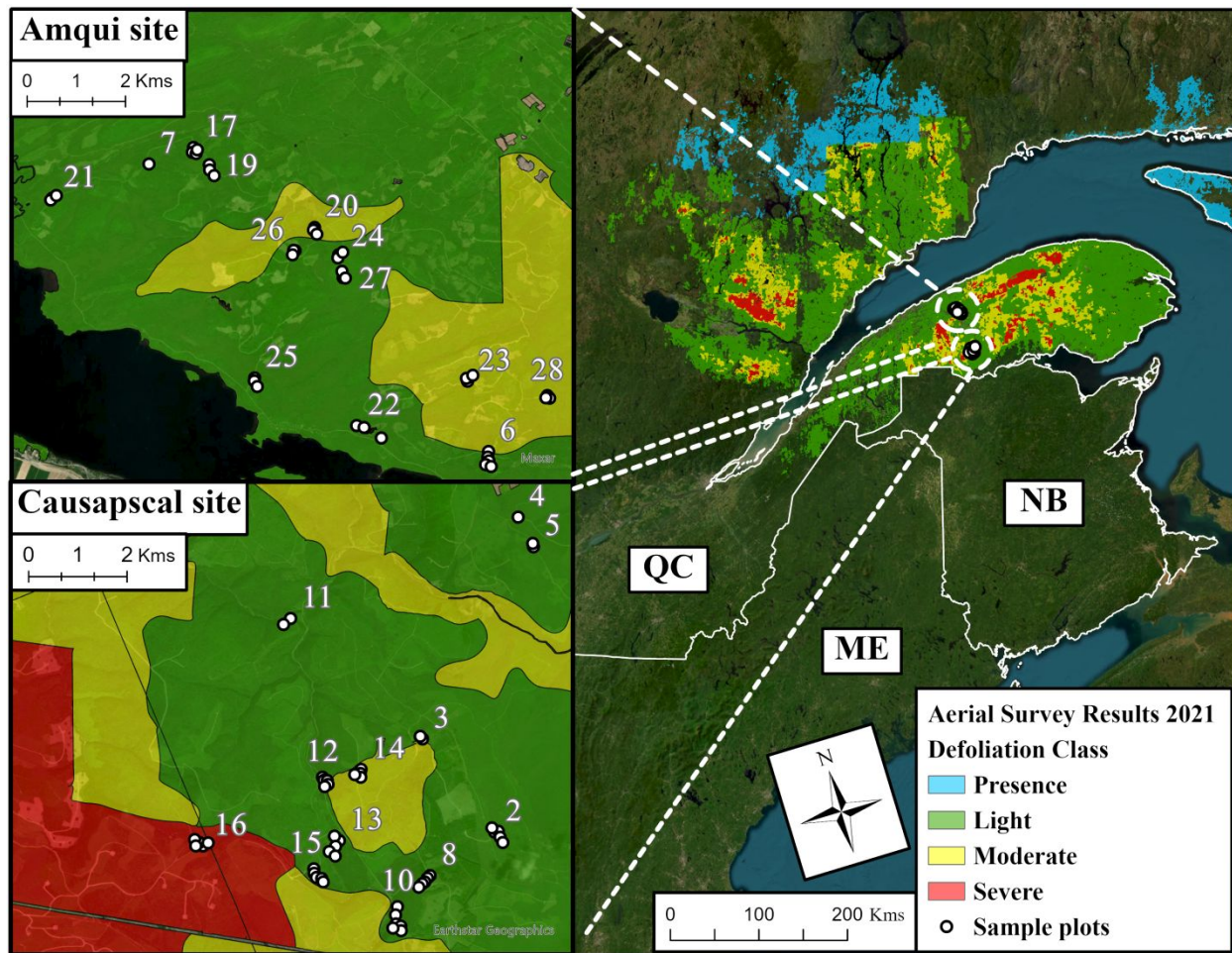
Model ~ Explanatory variables	AIC	BIC	Likelihood ratio test p-value	Marg. R ²	Cond. R ²	RMSE
LMM: Balsam fir cumulative defoliation						
M1 ~ Sprayed	8810	8838		0.014	0.938	43.066
M2 ~ Sprayed + SuMinTp	8807	8840	0.021	0.020	0.933	42.945
M3 ~ Sprayed + SuMinTp + WiMaxTp	8802	8840	0.008	0.054	0.933	42.765
M4 ~ Sprayed + SuMinTp + WiMaxTp + SuMaxTp	8799	8841	0.024	0.110	0.928	42.667
M4a ~ Sprayed + SuMinTp + SuMaxTp + L2branch*HWba	8754	8806	<0.0001	0.066	0.929	41.306
M4b ~ <i>Sprayed + SuMinTp + SuMaxTp + HWba + WSba + L2branch*HWba + L2branch*WSba</i>	8712	8773	<0.0001	0.085	0.931	40.186
M5 ~ Sprayed + SuMinTp + WiMaxTp + SuMaxTp + Totalba	8797	8844	0.051	0.111	0.928	42.691
M6 ~ Sprayed + SuMinTp + WiMaxTp + SuMaxTp + Totalba + BFba	8795	8847	0.045	0.111	0.928	42.692
GLMM: Balsam fir current defoliation						
M1 ~ Sprayed	4430	4454		0.031	0.340	0.176
M2 ~ Sprayed + L2branch	4340	4368	<0.0001	0.053	0.281	0.173
M3 ~ Sprayed + L2branch + WiMaxTp	4242	4274	<0.0001	0.205	0.557	0.167
M4 ~ Sprayed + L2branch + WiMaxTp + SuMinTp	4223	4260	<0.0001	0.268	0.629	0.166
M5 ~ Sprayed + L2branch + WiMaxTp + SuMinTp + SuMaxTp	4214	4257	0.0012	0.227	0.604	0.165
M5a ~ Sprayed + L2branch + WiMaxTp + SuMinTp + SuMaxTp + Sprayed*SuMinTp	4141	4188	<0.0001	0.209	0.582	0.161
M5b ~ <i>Sprayed + L2branch + WiMaxTp + SuMinTp + SuMaxTp + HWba + Sprayed*SuMinTp + L2branch*HWba</i>	4100	4156	<0.0001	0.193	0.542	0.160
M6 ~ Sprayed + L2branch + WiMaxTp + SuMinTp + SuMaxTp + SpPrcep	4210	4257	0.014	0.230	0.575	0.165
M7 ~ Sprayed + L2branch + WiMaxTp + SuMinTp + SuMaxTp + SpPrcep + SpMaxTp	4207	4259	0.028	0.191	0.507	0.165

871 Note: See Table 1 for description of model explanatory variables. Models M4a and M4b (cumulative defoliation) and M5a and M5b
 872 (current defoliation) evaluated adding interactions among explanatory variables.

873 **Table 5.** Summaries for final generalized and linear mixed-effects models (GLMM, LMM)
 874 examining the effects of site, stand, and climate explanatory variables on balsam fir current and
 875 cumulative defoliation response variables. Coefficient mean estimates and log-odds (converted
 876 probabilities) are based on standardized z-scores along with standard errors and significance.

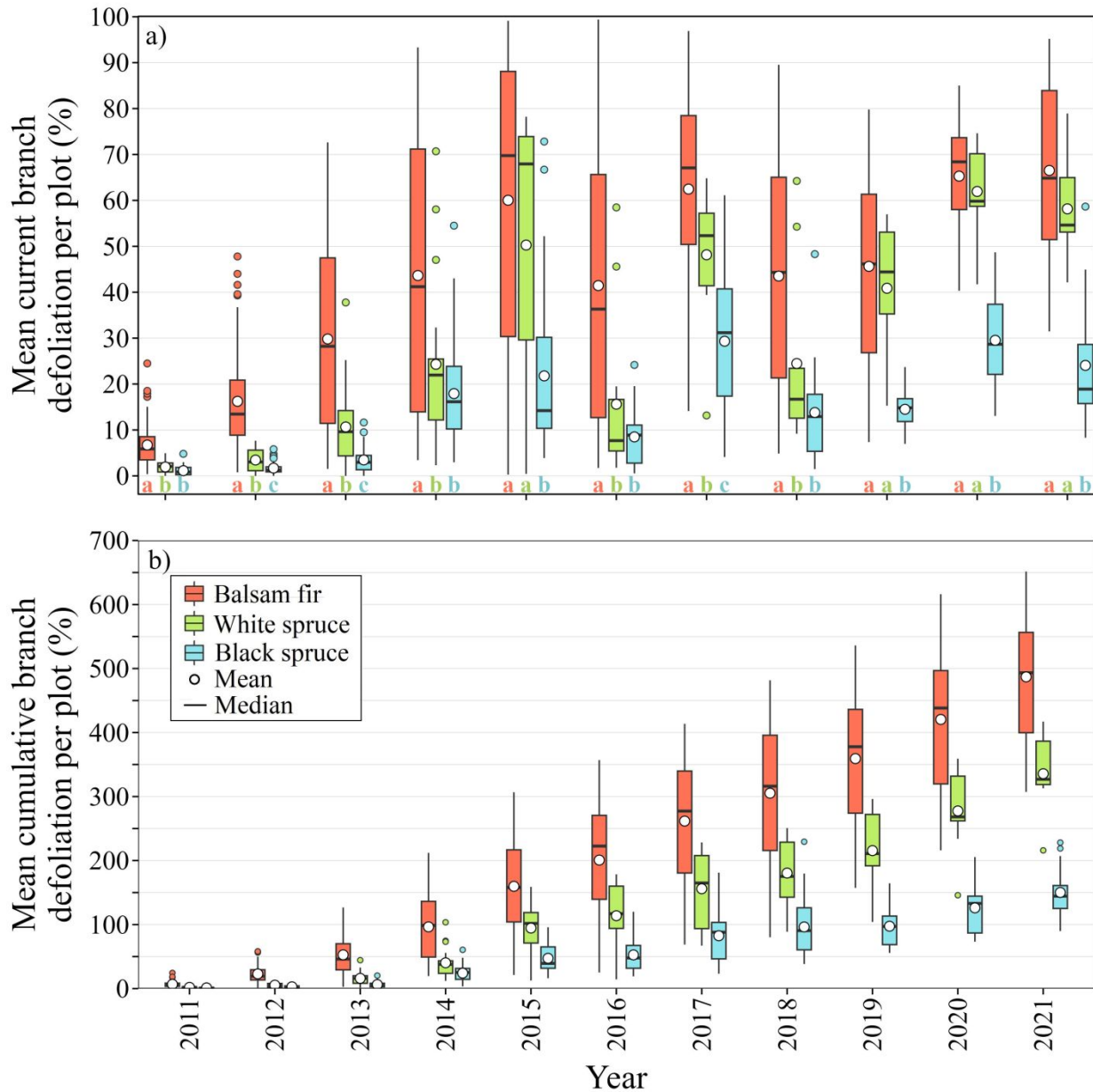
LMM: Balsam fir cumulative defoliation				
Fixed effect variables	Mean estimate	Std. error	t-value	p-value
Intercept	225.30	43.33	5.20	0.0003
Sprayed	-84.08	6.78	-12.40	<0.0001
SuMinTp	-19.33	5.81	-3.33	0.0009
SuMaxTp	29.83	11.34	2.63	0.0087
HWba	-17.09	7.55	-2.26	0.0282
WSba	-10.37	4.09	-2.54	0.0133
L2branch*HWba	-15.22	1.65	-9.25	<0.0001
L2branch*WSba	-8.53	1.32	-6.47	<0.0001
GLMM: Balsam fir current defoliation				
Fixed effect variables	Log-odds (probability)	Std. error	z-value	p-value
Intercept	-0.26 (0.43)	0.46	-0.57	0.57
Sprayed	-1.53 (-0.18)	0.11	-13.39	<0.0001
L2branch	0.48 (0.62)	0.04	11.41	<0.0001
SuMinTp	-0.17 (-0.46)	0.09	-1.99	0.0002
SuMaxTp	-0.61 (-0.35)	0.16	-3.92	<0.0001
WiMaxTp	1.02 (0.73)	0.13	7.62	<0.0001
HWba	-0.17 (0.46)	0.09	-1.96	0.05
Sprayed*SuMinTp	-0.94 (-0.28)	0.12	-8.05	<0.0001
L2branch*HWba	-0.16 (-0.46)	0.03	-6.28	<0.0001

877 **Note:** Abbreviations for fixed effect explanatory variables are listed in Table 1. Alternative
 878 model formulations used to select the final model based on lowest Bayesian Information
 879 Criterion (BIC) are presented in (Table 4). To convert the standardized model coefficients back
 880 to their original values/units, the formula is: $X = Z * (\sigma_x + \mu_x)$ where Z is the standardized value.



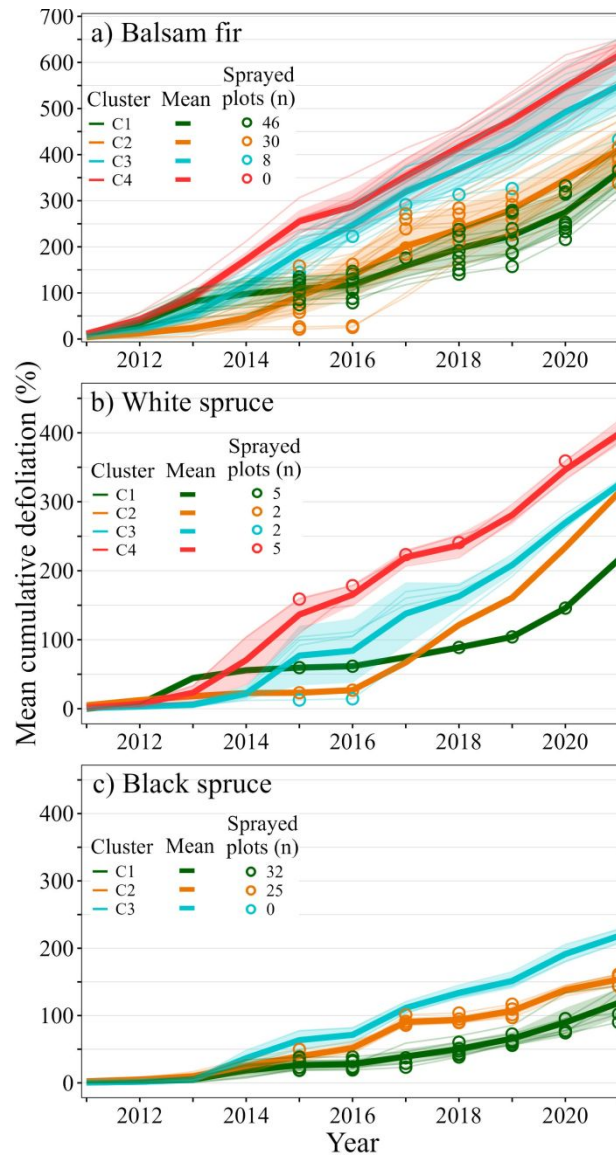
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882 **Fig. 1.** Study area for two sites containing the 87 sample plot locations (Amqui $n = 40$ and
 883 Causapscaal $n = 47$) overlaid on Québec government aerial surveys of spruce budworm
 884 defoliation classes from 2021 (MRNF, 2021). Groups of sample plots are labelled by their
 885 corresponding stand number; data on characteristics of each stand are included in Supplementary
 886 Table S1. Figure created using ArcGIS Pro 3.3.2. (ESRI, 2023) with base map satellite imagery
 887 (ESRI), aerial survey defoliation layer (MRNF, 2021), and overlaid points for plot locations in
 888 the WGS 1984 UTM 19N projection.



889

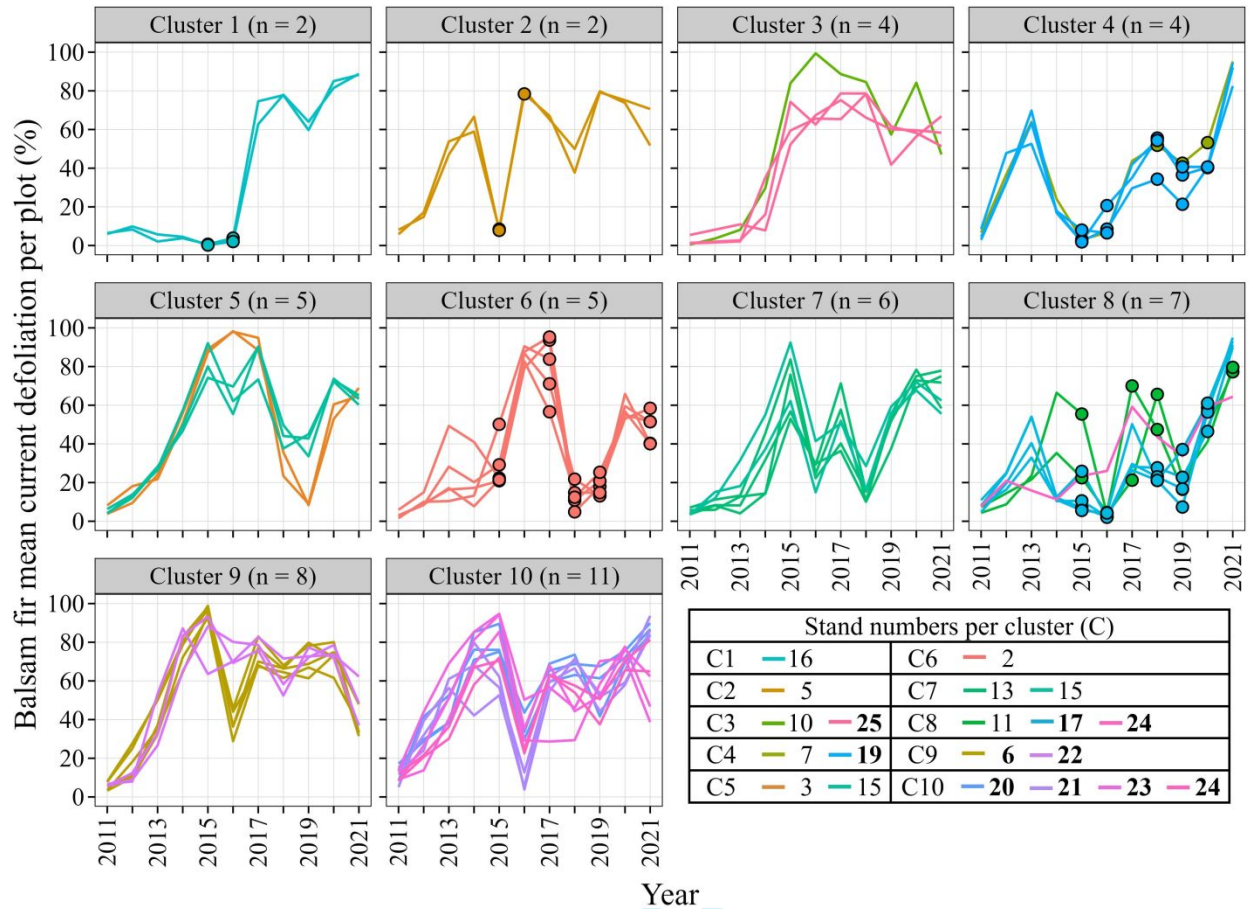
890 **Fig. 2.** Box plots showing (a) mean current defoliation and (b) mean cumulative defoliation by
 891 spruce budworm host tree species (balsam fir, white spruce, and black spruce) for 87 sample
 892 plots from 2011–2021. Colored letters at bottom of panel a) indicate significant differences of
 893 means among species per year using Games-Howell pairwise comparison test. Cumulative
 894 defoliation was calculated by summing mean current annual defoliation.



895

896

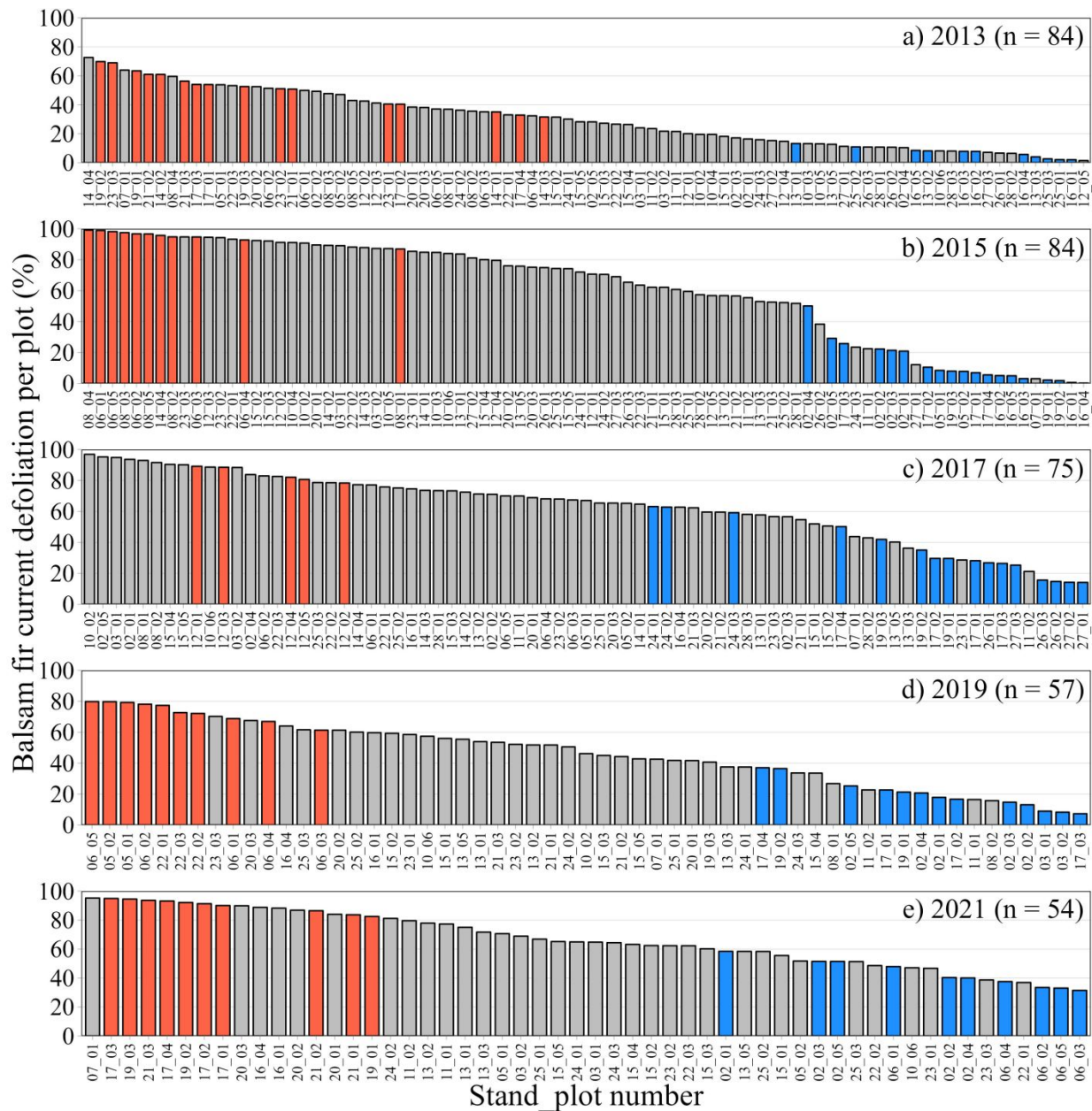
897 **Fig. 3.** Hierarchical cluster analysis results for cumulative defoliation patterns of plots from
 898 2011–2021 of (a) balsam fir, (b) white spruce, and (c) black spruce. Shaded bands represent ± 1
 899 standard deviation of the mean per cluster, fine lines show individual plots, and circles indicate
 900 years of annual aerial spraying of insecticide per plot.



901

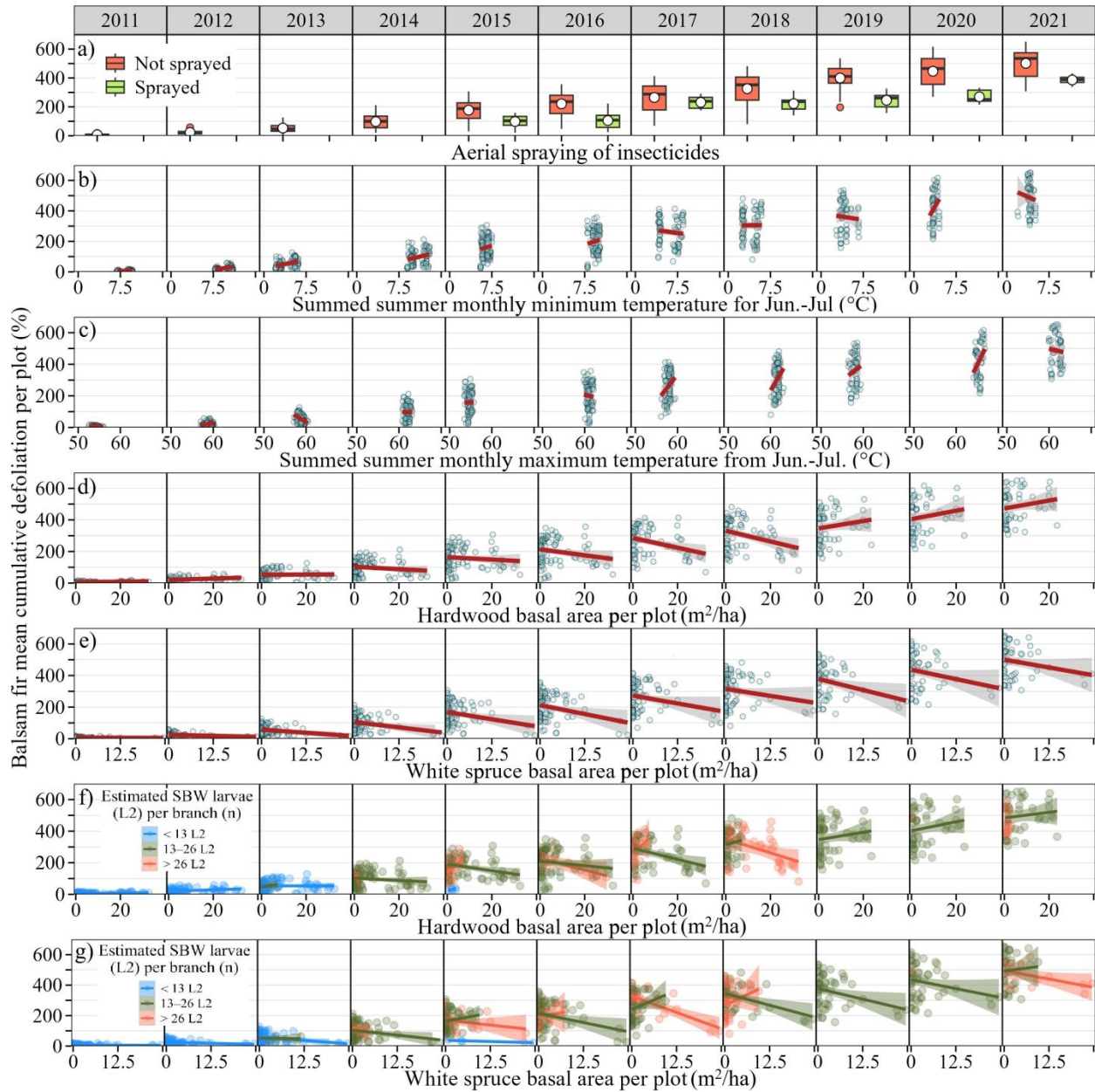
902

903 **Fig. 4.** Hierarchical cluster analysis results for current defoliation patterns from 2011–2021 of
 904 balsam fir. Each panel represents a cluster of plot defoliation patterns that are colored by stand
 905 number. Circles indicate years of annual aerial spraying of insecticide per plot and bold numbers
 906 in the legend indicate stands located in the Amqui site.



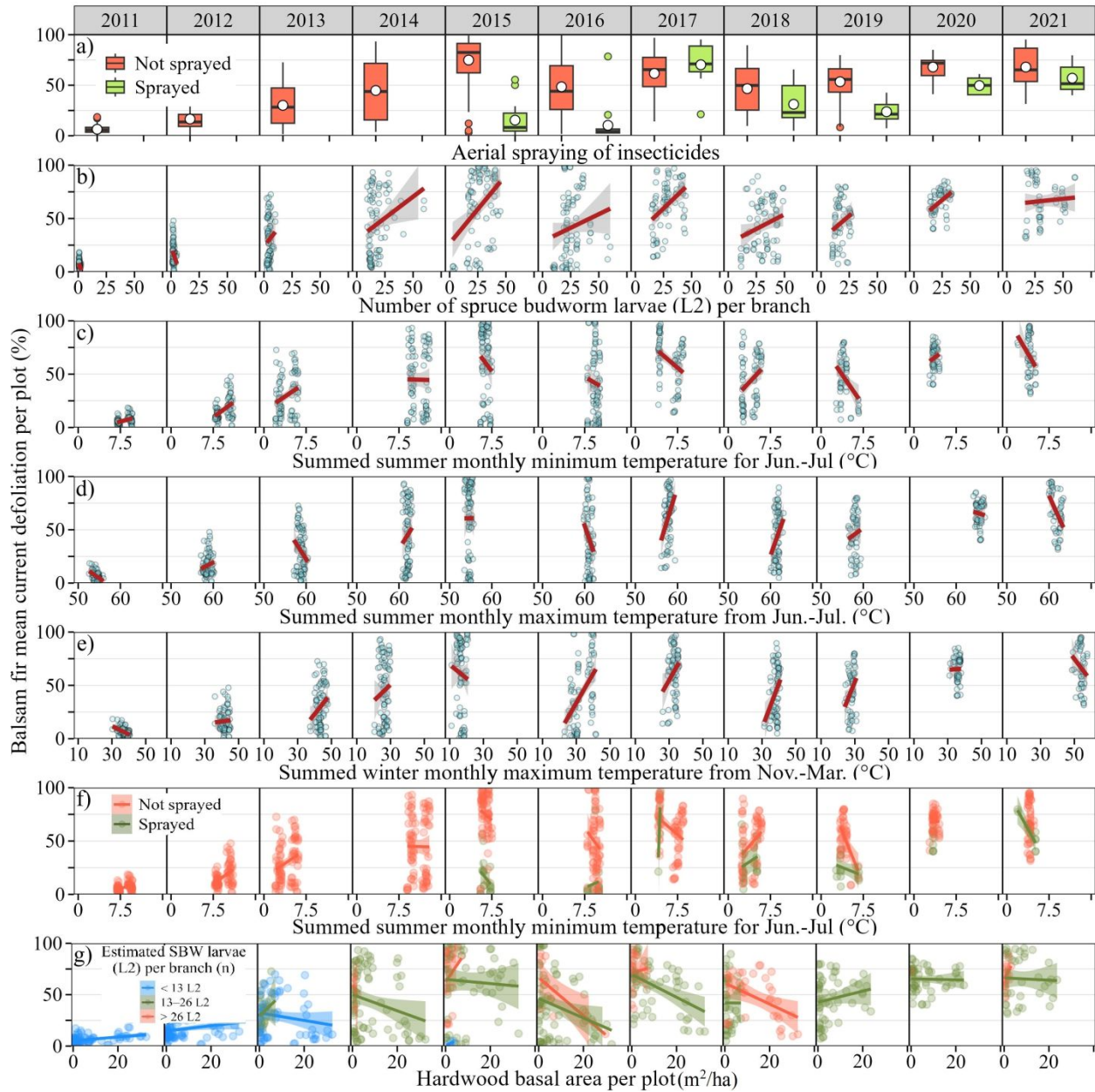
907

908 **Fig. 5.** Mean balsam fir current defoliation for plots, ordered from highest to lowest defoliation
 909 for every second year from 2013 to 2021. Plots showing significantly clustered within stand
 910 defoliation ($\alpha = 0.05$) are colored red ('hot spots' - highly defoliated plots surrounded by other
 911 highly defoliated plots) and blue ('cold spots' - lightly defoliated plots surrounded by other
 912 lightly defoliated plots), or grey (not significantly clustered), based on Getis-Ord G_i^* analysis.
 913 Number of plots per year declined due to timber harvesting.



914

915 **Fig. 6.** Fitted linear relationships between annual balsam fir mean cumulative defoliation per plot
 916 and significant explanatory variables: a) annual spraying of insecticides; b) summer monthly
 917 minimum temperature; c) summer monthly maximum temperature; d) hardwood basal area per
 918 plot; e) white spruce basal area per plot; f) interaction between hardwood basal area and SBW L2
 919 per branch; and g) interaction between white spruce basal area and SBW L2 per branch.



920

921 **Fig. 7.** Fitted linear relationships between annual balsam fir mean current defoliation per plot and
 922 significant explanatory variables: a) spraying of insecticides; b) SBW L2 populations per branch;
 923 c) summer monthly minimum temperature; d) summer monthly maximum temperature; e) winter
 924 monthly maximum temperature; f) interaction between spraying insecticides and summer
 925 cumulative degree days; and g) interaction between hardwood basal area and SBW L2.