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1	Using Time Series of MODIS Land Surface Phenology to Model Temperature and
2	Photoperiod Controls on Spring Greenup in North American Deciduous Forests
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12	

13 Abstract

The timing of leaf emergence in temperate and boreal forests is changing, which has 14 profound implications for a wide array of ecosystem processes and services. Spring phenology 15 16 models, which have been widely used to predict the timing of leaf emergence, generally assume 17 that a combination of photoperiod and thermal forcing control when leaves emerge. However, the exact nature and magnitude of how photoperiod and temperature individually and jointly 18 19 control leaf emergence is the subject of ongoing debate. Here we use a continuous development 20 model in combination with time series of land surface phenology measurements from MODIS to quantify the relative importance of photoperiod and thermal forcing in controlling the timing of 21 22 canopy greenup in eastern temperate and boreal forests of North America. The model accurately 23 predicts biogeographic and interannual variation in the timing of greenup across the study region (median RMSE = 4.6 days, median bias = 0.30 days). Results reveal strong biogeographic 24 variation in the period prior to greenup when temperature and photoperiod influence greenup that 25 26 covaries with the importance of photoperiod versus thermal controls. Photoperiod control on leaf 27 emergence is dominant in warmer climates, but exerts only modest influence on the timing of 28 leaf emergence in colder climates. Results from models estimated using ground-based 29 observations of cloned lilac are consistent with those from remote sensing, which supports the 30 realism of remote sensing-based models. Overall, results from this study suggest that apparent 31 changes in the sensitivity of trees to temperature are modest and reflect a trade-off between 32 decreased sensitivity to temperature and increased photoperiod control, and identify a transition 33 in the relative importance of temperature versus photoperiod near the 10 °C isotherm in mean 34 annual temperature. This suggests that the timing of leaf emergence will continue to move earlier

- as the climate warms, and that the magnitude of change will be more pronounced in colder
 regions with mean annual temperatures below 10 °C.
- 37 Keywords: spring phenology, climate change, deciduous forests, photoperiod, temperature
- 38 sensitivity, Bayesian, hierarchical modeling, MODIS, land surface phenology

40 **1. Introduction**

There is overwhelming evidence that leaf emergence is occurring earlier in temperate and 41 boreal forests (Menzel et al., 2006; Schwartz et al., 2006). However, a number of recent papers 42 have concluded that the sensitivity of leaf emergence to changes in temperature has decreased in 43 recent decades (Fu et al., 2015; Piao et al., 2017) and that the period when trees are sensitive to 44 45 thermal forcing is becoming shorter (Fu et al., 2019; Güsewell et al., 2017; Wenden et al., 2020). These results complicate interpretation of observed trends and exacerbate challenges involved in 46 47 forecasting how the phenology of trees will change in the future. These challenges are further complicated by fundamental issues in the way that the sensitivity of phenological events to 48 49 temperature is generally quantified (Keenan et al., 2019). Because changes in phenology impact 50 important ecosystem functions (Keenan et al., 2014; Richardson et al., 2013), understanding how changes in climate affect phenology is critical to forecasting how ecosystems will respond to 51 future climate change (Peñuelas et al., 2009; Piao et al., 2019). 52

53 To address this, a variety of recent studies have focused on improving understanding of 54 bioclimatic controls on plant phenology (Liu et al., 2017; Zohner et al., 2016). Results from both lab- and field-based experimental studies have provided insights (Montgomery et al., 2020; 55 Richardson et al., 2018a), but are limited by the fact that phenological behavior in controlled 56 laboratory- and field-based warming experiments differs from behavior observed in natural 57 ecosystems (Clark et al., 2014a; Wolkovich et al., 2012). Further, the manner in which 58 environmental conditions are perturbed in such experiments (e.g., 2 °C warming) is not 59 60 representative of climate changes expected in the future, which are predicted to occur gradually, but with large year-to-year variability (Schewe et al., 2019; Walther et al., 2002). These issues 61 are compounded by the fact that the geographic sampling of data sets used in these studies is 62

often limited and does not reflect the full biogeographic range of species examined (Richardson
et al., 2013). Hence, geographic variation in the relative importance of different climate drivers
on phenology, both within and across plant communities, is not well understood (Piao et al.,
2019).

67 One widely used strategy for investigating the response of plant phenology to climate change is to calibrate mechanistic models using weather data in combination with long-term records of 68 69 phenology collected on the ground (Basler, 2016; Fu et al., 2019) or from remote sensing (Liu et al., 2017; Melaas et al., 2018). In addition to thermal controls, photoperiod is widely assumed to 70 71 control the timing of leaf emergence by regulating the entrance of ecodormancy, triggering thermal forcing to stimulate bud swelling and leaf emergence (Chuine et al., 2016; Jackson, 2009; 72 Körner and Basler, 2010). Hence, many models include explicit representation of photoperiod 73 (e.g., Blümel and Chmielewski, 2012; Masle et al., 1989; Basler, 2016; Migliavacca et al., 2012). 74 75 To capture the role of thermal forcing, mechanistic models generally use aggregated bioclimatic variables such as growing degree days or winter chilling as their primary inputs. However, Clark 76 77 et al. (2014a) have suggested that the use of such aggregated quantities is problematic because 78 values for prescribed variables required by these models (e.g., start date of forcing accumulation) 79 are not identifiable.

In recent years, data-driven models based on state-space representations of phenological processes have been developed that overcome many of the weaknesses of both mechanistic and experimental approaches (e.g., Clark et al., 2014b; Qiu et al., 2020; Senf et al., 2017; Seyednasrollah et al., 2018). By modeling phenological dynamics directly from data, these models avoid issues arising from misspecification of functional relationships between forcing variables and processes that regulate phenological development (Clark et al., 2014b). Building on

this approach, here we use a data-driven spring onset model in combination with gridded weather 86 87 data and time series of ground-based and remotely sensed observations of spring greenup dates to explore biogeographic patterns in photoperiod and thermal controls on the timing of spring 88 89 greenup. Specifically, we use this model to : (1) quantify the relative importance of thermal forcing, photoperiod, and winter chilling in controlling spring greenup; (2) identify the pre-90 season period when plants are sensitive to bioclimatic controls; and (3) characterize how 91 92 covariance among thermal forcing, photoperiod, and the length of the pre-season period control 93 the biogeography of spring greenup in deciduous forests of eastern temperate and boreal North America. 94

95

96 **2. Methods**

97 2.1. Study Region

The study region includes the Northern Forests and Eastern Temperate Forest ecoregions included in level 1 of the US EPA Ecoregions of North America (Fig. A1). To distinguish deciduous forests from evergreen forests and other land cover types within the study area, the 500 m Collection 6 MODIS Land Cover Type product was used. This product provides annual land cover maps based on machine learning that are post-processed using a multi-temporal statespace modeling framework that reduces spurious land cover change introduced by classification uncertainty in individual years (Abercrombie and Friedl, 2016; Sulla-Menashe et al., 2019).

The continuous development spring onset model (Section 2.3) is estimated on an equal-area grid, where each grid cell is 4.67 km x 4.67 km (10×10 MODIS pixels; ~22 km²). In each grid cell, only pixels labeled as deciduous broadleaf or mixed forests throughout the entire study period from 2001-2017 were included in the analysis. To ensure analyses were based on grid 109 cells dominated by deciduous forest cover, we excluded model grid cells where the fraction of
110 MODIS pixels labeled as deciduous broadleaf or mixed forests was less than 50% (Fig. A1).

111

112 2.2. Spring Greenup and Meteorological Data

To identify the timing of springtime leaf emergence from 2001 to 2017, we used the 113 114 Collection 6 MODIS Land Cover Dynamics product (MCD12Q2; Gray et al., 2019). This product uses time series of the two-band Enhanced Vegetation Index (EVI2) to identify the 115 116 timing of six key phenophase transition dates during each growing season in each 500-m MODIS 117 pixel. Numerous studies have reported that this product provides a reliable measure of vegetation 118 phenology (Moon et al., 2019; Richardson et al., 2018b) and seasonal changes in ecological function and surface biophysical characteristics (Melaas et al., 2013; Moon et al., 2020). For this 119 120 analysis, we use the MCD12Q2 'greenup' metric, which is defined by the Land Cover Dynamics product as the day of year (DOY) during the greenup phase in spring when the EVI2 time series 121 122 at each pixel crosses 15% of its seasonal amplitude (Gray et al., 2019).

123 To provide the meteorological data required for model estimation, we used the Version 3 124 Daymet dataset for North America (Thornton et al., 2017) (https://daymet.ornl.gov). This data set uses digital elevation data in association with a land-water mask and meteorological 125 observations collected at ground-based stations to create gridded time series of surface 126 127 meteorological fields at daily time step and 1 km spatial resolution for the period 1980 to present. 128 For this work, we used daily maximum and minimum 2-m air temperatures from 2000 to 2017 129 along with daylength, resampled to 500 m and co-registered to the MODIS data over all grid 130 cells included in our analysis.

132 **2.3. Continuous Development Spring Onset Model**

133 To estimate the sensitivity of different climatological controls on springtime phenology, we developed a continuous development spring onset model (hereafter, CDSOM) based on a 134 hierarchical Bayesian framework that predicts the timing of springtime greenup using three 135 136 drivers: photoperiod, thermal forcing, and chilling units. The original form of this model was proposed by Clark et al. (2014b), who used the same general approach to show that because 137 conventional process-based phenology models (e.g., Hufkens et al., 2018.) aggregate daily air 138 139 temperature time series into cumulative sums or mean values for each year or season, they misrepresent how thermal forcing controls the timing of phenology. 140

Similar to Clark et al. (2014b), the CDSOM we use here tracks the continuous response of phenological development to variation in environmental controls at daily time step. To do this, the model uses a state-space framework that includes an unobservable latent state (h), which responds continuously to environmental controls and captures ecological responses to bioclimatic forcing:

$$h_{g,s,d+1} = h_{g,s,d} + \delta h_{g,s,d} \tag{1}$$

146 where $h_{g,s,d}$ is the latent state for grid cell g and sample (i.e., pixel) s on day d. In this framework, 147 $\delta h_{g,s,d}$ is the increment in h from day d to day d + 1, which is estimated using:

$$\delta h_{g,s,d} = \begin{cases} (X_{g,s,d} \times \beta_g)(1 - h_{g,s,d}/h_{max}), & \delta h_{g,s,d} \ge 0\\ 0, & \delta h_{g,s,d} < 0 \end{cases}$$
(2)

where $X_{g,s,d}$ is a matrix of predictor variables that includes the daily mean temperature $(T_{g,s,d})$, day-length (i.e., photoperiod; $L_{g,s,d}$), and chilling units (CU_{g,s}; defined below) on each day, and where daily mean temperature is computed as the average of daily maximum and minimum temperatures from Daymet in each 500 m MODIS pixel. β_g is a vector of estimated model coefficients for each grid cell (g), and h_{max} is the final state value of h, which is prescribed to be 100. Note that: (1) even though a linear formulation is used to describe the relationship between model predictors and coefficients, the model accommodates nonlinear responses in phenological responses to environmental controls using an asymptotic limit for the latent state (i.e., $h_{g,s,d}/$ h_{max}); and (2) the latent state increment is always non-negative.

157 To convert the continuous scale of the latent state (h) into a form that identifies discrete 158 phenological events (i.e., the timing of spring greenup onset), a logit transformation is used:

$$logit(P_{g,s,d}) = \kappa + \lambda \times h_{g,s,d}$$
(3)

159 where $P_{g,s,d}$ is the probability that the onset occurs at sample pixel *s* in grid *g* on day *d*, and κ and 160 λ are the intercept and slope of the transformation, respectively. Because greenup onset is 161 defined to be a discrete event, $P_{g,s,d}$ follows a Bernoulli distribution:

$$Y_{g,s,d} \sim Bernoulli(P_{g,s,d}) \tag{4}$$

162 where $Y_{g,s,d}$ indicates whether or not greenup onset has occurred for sample s in grid g on day d.

163 Following convention, chilling units $(CU_{q,s})$ were defined as:

$$CU_{g,s} = \sum_{d=c_{g,0}}^{c_{g,s}} I(T_{g,s,d} < T_b)$$
(5)

Hence, $CU_{g,s}$ is defined as the number of days below prescribed threshold T_b during the period after the onset of dormancy until an unobserved date $c_{g,s}$ when the chilling requirement is satisfied. Previous studies have suggested that boreal and temperate tree species respond to air 167 temperatures ranging from -5 to 10 °C as a threshold for chilling requirements (Hänninen et al., 168 2019). Here we used 0 °C because the study area covers a large range of climate conditions. 169 Further, and more importantly, sensitivity analyses revealed that model results were not sensitive 170 to variation in T_b (not shown), which is supported by results indicating that chilling control on 171 the timing of greenup is minor (see Results).

172

173 2.4. CDSOM Estimation

174 As we described above, the CDSOM was estimated using a regular grid, with each grid 175 cell composed of 100 MODIS pixels. We excluded all pixels with more than one land cover type label between 2001 and 2017 (i.e., that nominally experienced change) and excluded all cells that 176 177 were composed of less than 50% deciduous or mixed forests. Because the CDSOM is 178 computationally expensive, we used a two-stage sampling approach to estimate the model for 179 randomly selected grid cells in each of the 13 MODIS tiles that intersect the study region. In the 180 first stage, we randomly sampled grid cells within each MODIS tile that met the criteria listed 181 above. If less than 300 valid grid cells were available within a tile, we included all valid grid cells. If more than 300 grid cells were available in a tile, we randomly selected a sample of 300 182 183 cells. In the second stage, we randomly selected MODIS pixels located in each grid cell across 184 time. To minimize the impact of spatial and temporal correlation, we used a sub-sample of 100 185 pixel-years (i.e., 100 unique greenup dates randomly selected across 17 years) to estimate a unique model for each cell. Each sample was selected from a total pool of between 850-1700 186 187 sample points (i.e., 50-100 pixels per year in each grid across 17 years).

For each year, December 1st of the previous year and DOY 250 (~Sept. 7) of the current year
were used as the start and end dates of latent state development, respectively. Posterior sampling

190 was performed using the "R2jags" package in R (Su and Yajima, 2015), with 10,000 iterations 191 and 3,000 burn-in periods. As a final step, to reduce noise in our results, we excluded grid cells 192 where estimated model coefficients were outside 95% of the range of estimated model 193 coefficients across all grid cells. This yielded a final data set consisting of 1,685 grid cells with 194 valid results.

195

196 Model results from a representative grid cell are shown in Fig. 1. Overall, predicted onset 197 dates are well aligned with observed onset dates at this grid cell, with a root-mean-square error 198 (RMSE) of 3.7 days across the time series (Fig. 1a). Because the input forcing data are normalized prior to model estimation (i.e., having a mean of 0 and a standard deviation of 1 for 199 each of the input variables in each grid g and sample s), the posterior distributions for each 200 201 model coefficient, which reflect the dependence of phenological development on each input 202 variable, show differences that are independent of the magnitude or units of each input variable (Fig. 1b). Time series of the latent state generated by the model (Fig. 1c) provide information 203 regarding the timing and duration of the pre-season period prior to greenup onset. This period 204 has been previously described as "as the most temperature-sensitive period preceding the 205 phenological event" (Güsewell et al., 2017) or "the period before leaf unfolding for which the 206 partial correlation coefficient between leaf unfolding and air temperature is highest" (Fu et al., 207 2015). Here we define this period as corresponding to the time interval when phenological 208 209 development is affected by bioclimatic forcing, and we use the CDSOM to identify the "preseason period" as starting on the DOY when the latent state variable (h) starts to increase and 210 ending on the DOY when greenup onset occurs (i.e., the period indicated by the arrow in Fig. 1c). 211



Fig. 1. Model results for a randomly selected grid cell. (a) Relationship between the greenup 213 onset dates from MODIS and onset dates estimated by the model. (b) The distribution of model 214 coefficients for each control variable (i.e., the relative dependence on each climate control; 215 Therm.: thermal forcing; Photo.: photoperiod; Chill.: chilling units). (c) Time series of the latent 216 217 state (red line) and the length of the pre-season (identified by the horizontal arrow). In panel (a), 218 each dot (total n = 100) represents an individual pixel-year sampled from the grid cell comprised 219 of 10 by 10 MODIS pixels across 17 years of the study period (i.e., 100 out of the total 1,700 pixel-years). 220

222 **2.5.** Quantifying the Relative Importance of Bioclimatic Forcing Variables

223 To address our goal of quantifying the relative importance (and geographic variation thereof) among bioclimatic controls on the timing of springtime phenology, we compute a normalized 224 225 index with values that range from -1 to +1 that captures this effect. Because each of the input variables in each grid g and sample s have been normalized to have a mean of 0 and a standard 226 deviation of 1, model coefficients can be directly compared to assess the relative importance of 227 228 each control variable. To quantify this, we calculated the relative importance (RI) of each control 229 variable relative to each other variable using a normalized index computed from CDSOM model 230 results. For example, to compute the relative importance of photoperiod versus thermal forcing 231 in any given grid cell, we computed:

$$RI = \frac{\beta_T - \beta_L}{\beta_T + \beta_L} \tag{6}$$

where β_T and β_L are the average model coefficients for thermal forcing and photoperiod (respectively) during the pre-season period, which are estimated for each grid cell by the CDSOM.

235

236 2.6. CDSOM Assessment and Comparison with Conventional Phenology Models

To provide a baseline comparison against previously developed and widely used springtime phenology models (hereafter, the 'conventional models'), we compared results from the CDSOM with four widely used process-based phenology models included in the "phenor" package in R (Hufkens et al., 2018). Specifically, we compared our results against the thermal time (TT) model, the photo-thermal time (PTT) model, the exponential photo-thermal time model (M1), and the alternating (AT) model, as described by Hufkens et al. (2018). These models are fundamentally different from the CDSOM in that they assume a linear relationship between spring thermal forcing and the rate of phenological development, and that spring onset occurs when accumulated forcing (after a prescribed start date) reaches a critical threshold (F^*). The TT model relies only on thermal forcing (daily air temperature in each MODIS pixel, $T_{g,s,d}$) with no additional inputs. In this model, the state of forcing (S_f) increases each day until F^* is reached, when leaves emerge (Chuine et al., 1999; Hunter and Lechowicz, 1992):

$$R(T_{g,s,d}) = \begin{cases} 0 & \text{for } T_{g,s,d} \le T_b \\ T_{g,s,d} - T_b & \text{for } T_{g,s,d} \ge T_b \end{cases}$$
(6)

$$S_f = \sum_{t_0}^t R(T_{g,s,d}) \tag{7}$$

where t_0 is the starte date. For consistency with the CDSOM, we set t_0 and T_b to December 1st and 0 °C, respectively. The PTT model includes day-length (i.e., photoperiod; $L_{g,s,d}$) as an additional factor that regulates the rate of thermal forcing (Črepinšek et al., 2006; Masle et al., 1989):

$$S_f = \sum_{t_0}^t R(T_{g,s,d}) \times \frac{L_{g,s,d}}{24}$$
(8)

The exponential M1 model also includes photoperiod, but treats the relationship between photoperiod and S_f as an exponential (Blümel and Chmielewski, 2012):

$$S_f = \sum_{t_0}^t R(T_{g,s,d}) \times \left(\frac{L_{g,s,d}}{24}\right)^k \tag{9}$$

where *k* is an empirically estimated constant. Finally, the AT model includes the number of days when the daily mean temperature falls below T_b (i.e., the number of chilling days; NCD), and treats NCD as an exponential function that reduces the thermal forcing accumulation required for
spring onset to occur (Cannell and Smith, 1983):

$$F^* = a + b \times \exp[c \times \text{NCD}(t)] \tag{10}$$

where *a*, *b*, and *c* are empirically estimated constants, and NCD(t) is defined as the number of chilling days since December 1st. A table summarizing the variables and main characteristics of the CDSOM and conventional models is provided as an appendix (Table A1).

262 For this analysis, we assessed model performance for both the CDSOM and the conventional 263 models in two ways. First, we assessed results from model-based predictions for the timing of spring greenup based on all available years (from 2001 to 2017). Second, to provide a more 264 robust assessment of model performance, we held out two years (2010 and 2012) with 265 anomalously warm springs in much of the study region (Friedl et al., 2014), and evaluated model 266 267 performance for each of these years. In this way, we were able to assess not only how well the 268 models performed under average conditions, but also how well they performed under unusual springtime weather conditions that were not represented in the data used to estimate the models. 269

270

271 2.7. CDSOM Estimation Using Ground-Based Observations

As a final element of our analysis, to complement model results based on remotely sensed greenup dates and to provide an independent basis for assessing the realism and robustness of our results, we estimated the CDSOM using time series of leaf unfolding dates for cloned lilac (*Syringa x chinensis* 'Red Rothomagensis') (Rosemartin et al., 2015). By applying the model to data from cloned plants, genetic variability is eliminated, and which allows us to investigate how differences in the timing of leaf unfolding between different individuals are caused by

differences in local environmental controls. Unlike our approach using MODIS spring greenup 278 279 dates, the model is estimated by pooling site-years across the region because the number of lilac leaf-out dates for each location is too small to accurately estimate models for each site. The 280 281 dataset includes 254 leaf unfolding dates from 60 locations across the study region, spanning the period from 2001 to 2008 (Fig. A1). For reasons we explain below, we stratified the dataset into 282 'warm' versus 'cold' sites based on whether the mean annual temperature at each site is above or 283 284 below 10 °C. Based on this stratification, the model was applied to 182 and 72 leaf unfolding 285 dates for the colder and warmer regions, respectively.

286

287 **3. Results**

288 The CDSOM accurately predicts biogeographic and interannual variation in the timing of springtime greenup across the study region. The median RMSE between predicted and observed 289 spring greenup dates was 4.6 days (Fig. 2), which is roughly equivalent to the uncertainty in 290 spring greenup dates estimated from MODIS (Moon et al., 2019). Inspection of results from the 291 292 conventional spring onset models show that median RMSEs were ~20% larger (~5.5 days vs. 4.6 days) relative to those obtained from the CDSOM (Fig. 2b). Further, RMSEs for years with 293 294 anomalous springs (2010 and 2012) were unchanged for the CDSOM, but increased by roughly 2 295 days for conventional models when 2010 and 2012 were excluded during model estimation (Fig. 3). For completeness, Fig. A2 shows the relationship between anomalies in MODIS greenup 296 dates and anomalies in predicted onset set dates, and demonstrates that the CDSOM outperforms 297 298 the conventional models in capturing year-to-year variations in spring onset dates. These results suggest that the CDSOM not only provides more accurate predictions of greenup relative to 299 predictions from conventional phenology models, but that the CDSOM more effectively captures 300

301 the impact of geographic and year-to-year variation in bioclimatic controls. More generally, the 302 accuracy of CDSOM results indicates that the model realistically captures the nature and 303 magnitude of ecophysiological responses to interannual and biogeographic variation in climate 304 controls that regulate the timing of greenup.

305



306

Fig. 2. Continuous Development Spring Onset Model (CDSOM) performance. (a) Geographic
variation in model root-mean-square error (RMSE) between greenup onset dates observed from
MODIS and onset dates predicted by the CDSOM model. (b) Boxplots showing the distribution
of RMSEs for the CDSOM model and four widely used conventional spring greenup models.
M1: The exponential photo-thermal time model; PTT: The photo-thermal time model; TT: The
thermal time model; AT: The alternating model. In panel (b), boxplots are presented in
increasing order of magnitude with respect to mean RMSE.

314



Fig. 3. RMSE results across models for anomalous years. (a) Boxplots of RMSEs for each model
for 2010 and 2012. (b) Boxplots showing increase in RMSEs for model predictions for all years
versus anomalous years (i.e., RMSEs for 2010 and 2012 – RMSEs for 2001-2017) at each grid
cell. CDSOM: continuous development spring onset model; M1: The exponential photo-thermal
time model; PTT: The photo-thermal time model; TT: The thermal time model; AT: The
alternating model. Boxplots are presented in increasing order of magnitude with respect to mean
RMSE.



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Fig. 4. Geographic variation in the dependence of spring greenup onset date to: (a) thermal forcing, (b) photoperiod, and (c) chilling units. In panel (d), boxplots show the distribution of model coefficients for each control variable during the pre-season period prior to leaf emergence in Northern Forests (blue) versus Eastern Temperate Forests (red). Differences between the means in both cases are statistically significant (p < 0.001).

333 The dependence of spring greenup on thermal forcing estimated by the CDSOM is higher in 334 Northern Forests than in Eastern Temperate forests (Fig. 4), but overall differences, while 335 statistically significant, are modest. In contrast, dependence on photoperiod control exhibits 336 systematic geographic variation across the study domain, with large differences between each ecoregion. Eastern Temperate Forests, which are warmer, show substantially higher dependence 337 on photoperiod relative to the Northern Forests ecoregion, which is much cooler (Fig. 4b and 4d). 338 339 This difference is especially pronounced in Eastern Canada where dependence on photoperiod is 340 low, versus the Southern United States, where photoperiod dependence is high. Dependence of spring onset on chilling units is uniformly low throughout the study region, which indicates that 341 342 the influence of chilling control, relative to photoperiod and thermal forcing, is effectively 343 negligible (Fig. 4c and 4d).

344 Geographic patterns in the RI of photoperiod versus thermal forcing indicates that 345 photoperiod exerts proportionally more control on the timing of spring greenup in warmer regions, while thermal forcing exerts proportionally more control in colder regions (Fig. 5a). By 346 plotting the *RI* in climate space (i.e., as a function of mean annual temperature and precipitation) 347 348 (Fig. 5b), the pattern becomes even more clear. In regions where mean annual temperature is above ~10 °C, photoperiod exerts stronger control on the timing of spring greenup than thermal 349 forcing. Conversely, in regions where mean annual temperature is less than ~10 °C, thermal 350 forcing is more important. RI values near the 10 °C isotherm in mean annual temperature are 351 generally close to zero, indicating equal influence of thermal forcing and photoperiod (plotted as 352 353 purple points in Fig. 5). These results suggest that the 10 °C isotherm in mean annual temperature identifies a transition zone between regions where thermal forcing versus 354 355 photoperiod is more dominant.



Fig. 5. Relative importance (RI) of thermal forcing versus photoperiod. Circles in red and blue show locations where thermal forcing and photoperiod, respectively, exert stronger control on the timing of spring greenup; purple circles identify locations where the magnitude of thermal forcing and photoperiod are roughly equivalent. The size of each circle is proportional to the magnitude of RI in each cell.

Results from applying CDSOM to ground-based observations of leaf unfolding dates for cloned lilac reveal that even though the individual lilac plants are genetically identical, the relative dependence of leaf unfolding dates on thermal forcing versus photoperiod depends on local bioclimatic conditions (Fig. 6). Consistent with previous studies (Basler and Körner, 2012; Schwartz et al., 2006), model coefficients and *RI* values indicate that leaf unfolding in cloned lilac depends more strongly on thermal forcing than on photoperiod, irrespective of location. However, thermal control is stronger in colder regions and *RI* values are significantly smaller

(i.e., thermal control is less dominant) in warm sites than in cold sites. In addition, comparison of cloned lilac data against greenup dates from MODIS for the same location show that MODIS greenup dates are biased late relative to lilac unfolding dates (Fig. A3), especially in warmer areas with earlier greenup dates, which supports the conclusion that lilacs are sensitive to temperature.

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Fig. 6. Dependence of cloned lilac leaf unfolding date on thermal forcing and photoperiod, and relative importance (*RI*). 254 total leaf unfolding dates from cloned lilac were divided into two groups based on mean annual temperature (≤ 10 °C, n = 182; > 10 °C, n = 72). The left panel plots the mean dependence of leaf unfolding on thermal forcing and photoperiod estimated by the CDSOM. The right panel plots the mean *RI* in each group. Positive *RI* indicates stronger control by thermal forcing relative to photoperiod. Vertical lines show ±1 standard deviation.

385 Finally, results from the CDSOM reveal patterns of covariance among pre-season period 386 length, photoperiod, and thermal forcing that jointly control the timing of greenup that are not 387 captured in conventional models. In particular, geographic variation in the pre-season period is strongly and negatively correlated with geographic variation in the relative importance of 388 photoperiod on spring greenup. Fig. 7b shows that this relationship follows a power law, where 389 photoperiod control decreases ($R^2 = 0.70$, p < 0.001) as the length of the pre-season period 390 391 increases. Fig. 7b also reveals modest heteroscedasticity in the relationship between pre-season 392 period length and photoperiod control, which reflects the fact that spring greenup in locations 393 with cooler mean annual temperatures and longer pre-seasons have lower dependence on 394 photoperiod and higher dependence on thermal forcing (Fig. 5). In contrast, the relationship between pre-season period and dependence on thermal forcing is statistically significant, but 395 much weaker ($R^2 = 0.13$; Fig. A4). 396

397



Fig. 7. Variation in pre-season period and the relationship between greenup dependence on photoperiod and length of pre-season period. (a) Geographic pattern in pre-season period, and (b) log-log relationship between the dependence of greenup on photoperiod and the length of the pre-season period.

403

404 **4. Discussion**

We assessed the relative importance of photoperiod, chilling, and thermal forcing in 405 controlling the timing of leaf emergence in Eastern Temperate and Boreal Forest ecoregions of 406 407 North America. To do this, we used a hierarchical Bayesian model in combination with time series of land surface phenology measurements from remote sensing. The former provides a 408 409 data-driven framework for investigating how different bioclimatic controls influence the timing 410 of leaf emergence (Clark et al., 2014b; Seyednasrollah et al., 2020); the latter provides a robust 411 and repeatable means of measuring and monitoring phenological dynamics over large areas 412 (Bolton et al., 2020; Zhang et al., 2018).

413 The core hypotheses that motivate this research include two main elements. First, the ecophysiological processes that control leaf emergence respond continuously to variation in 414 415 environmental controls throughout pre-season period prior to greenup in a manner that is not 416 represented in conventional models (Clark et al., 2014b). Second, rather than simply acting as a 417 cue for entering ecodormancy, photoperiod exerts continuous control on the timing of greenup 418 during the pre-season period. The results presented in this study suggest that both hypotheses are 419 supported. The preseason period, which corresponds to the period when the CDSOM latent state 420 variable (h) responds to bioclimatic forcing (Fig. 1c), ranges from roughly 2-12 weeks over the study domain (Fig. 7a). Throughout this period, changes in h reflect the net effect of daily 421

422 changes thermal and photoperiod controls. By estimating the model in a spatially explicit fashion 423 over a large geographic and climatic range, CDSOM results provide an empirical basis for 424 quantifying not only how thermal forcing and photoperiod individually and jointly influence the 425 timing of greenup, but more generally, how the length of the preseason period and relative 426 importance of photoperiod versus thermal forcing vary over the study domain.

427 Conventional models calibrated using long-term observations of phenological events such as 428 those used in this study have been widely used to simulate and forecast phenological events for 429 decades (Chuine and Régnière, 2017). Like the CDSOM, these models generally use air temperature, photoperiod, and chilling units in different configurations and combinations to 430 parameterize the response of plants to bioclimatic controls and predict the timing of phenophase 431 transitions (Basler, 2016; Hufkens et al., 2018). However, as we described previously, Clark et al. 432 433 (2014a, 2014b) argue that most conventional phenology models are fundamentally limited 434 because: (1) they aggregate measurements with substantial day-to-day variability over periods of weeks-to-months into single parameters and therefore do not capture how short-term variability 435 436 in control variables influences the timing of leaf emergence; (2) they rely on parameters that are 437 not identifiable; and (3) they do not account for uncertainty in model predictors or leaf emergence data. As a solution, Hänninen et al. (2019) argue that carefully designed factorial 438 439 experiments provide the most robust basis for improving understanding of processes that control 440 leaf emergence, and hence, for developing and testing process-based models. However, implementing such studies is difficult and expensive, and collecting sufficient sample data to 441 support robust and generalizable models is generally not possible. Reflecting these challenges, 442 results from a meta-analysis of warming studies showed that phenological changes observed in 443

such experiments do not replicate the magnitude of phenological responses to natural variation inair temperature observed in natural systems (Wolkovich et al., 2012).

Data-driven models like the CDSOM are not a panacea, but they do resolve several of the 446 issues discussed above. In addition to addressing the three limitations identified by Clark et al. 447 448 (2014a, 2014b), functional relationships among control variables in CDSOM are entirely 449 estimated from data. Hence the CDSOM avoids issues related to misspecification of functional 450 relationships that are inherent to conventional models. Further, by exploiting time series of 451 remote sensing observations collected over large areas that span nearly two decades, the CDSOM results presented here capture and reflect a much broader range of climate regimes and 452 climate variability than is generally possible using designed experiments. Indeed, we posit that 453 natural variability captured through interannual variability in climate over large geographic 454 455 scales provides an important and useful strategy for characterizing and understanding the 456 sensitivity of plant phenology to climate change (Friedl et al., 2014).

Moreover, and perhaps most importantly, while the patterns presented in Figs. 4-7 are 457 superficially consistent with results from previous studies suggesting that the timing of spring 458 459 greenup in deciduous forests has become less sensitive to thermal forcing and that the so-called 'temperature sensitive period' of temperate and boreal trees is changing (Fu et al., 2019, 2015; 460 Piao et al., 2017). We suggest that this inference may be spurious. Specifically, results from the 461 CDSOM show that thermal forcing control on the timing of greenup is heterogeneous and 462 463 exhibits weak covariance with pre-season period. Hence, apparent decreases in temperature 464 sensitivity actually reflect shorter pre-season periods with increased photoperiod control (Keenan et al., 2019). Stated another way, as the climate warms, higher temperatures tend to increase the 465 relative importance of photoperiod, while dependence on temperature has remained relatively 466

467 constant. Further, in regions where mean annual temperature is below ~10 °C, which 468 encompasses a significant proportion of the temperate zone and all of the boreal zone, 469 photoperiod control is modest and thermal forcing is clearly the dominant control. Indeed, our 470 results suggest that the biogeographic range in which the relative importance of photoperiod 471 control is increasing is restricted to locations with mean annual temperatures between ~8-10 °C, 472 and hence, is relatively narrow.

The simplest explanation for why photoperiod control varies geographically is provided by 473 474 the "law of the minimum", which states that plant growth is controlled by the scarcest resource rather than by the total resources available (Liebig et al., 1841). Our results are, to a first order, 475 476 consistent with this law. In cold regions (i.e., identified here as regions where mean annual 477 temperature is less than ~10 °C; Fig. 5), temperature is the primary limiting factor that controls the timing of greenup. In warmer regions where temperature is less limiting, light (or moisture) 478 becomes the primary limiting resource. Invoking a similar argument, Park et al. (2019) suggest 479 480 that extensive areas of high-latitude ecosystems that were previously constrained by temperature 481 are becoming more sensitive to photoperiod. Further, the results from our study are consistent 482 with recent experimental results from Zohner et al. (2016), who concluded that springtime phenology in deciduous trees at lower latitudes tended to depend more strongly on photoperiod, 483 484 while species at high latitudes leafed out independent of photoperiod. Hence our results are consistent with both long-established and more recent ecological literature. 485

Lastly, it is important to note several limitations of the current study. First, rather than modeling the role of chilling in controlling spring greenup using continuous (i.e., daily) forcing (Hänninen et al., 2019; Murray et al., 1989), the CDSOM uses chilling units, which provide an accumulated measure chilling requirements. This suggests that the role of the chilling units may 490 not be fully accounted for in this study, and may explain the relatively minor role of chilling 491 units in predicting the timing of spring greenup that we observed in this study (Fig. 4c and d) 492 (c.f., Heide and Prestrud, 2005; Laube et al., 2014). Second, to capture the effect of thermal forcing, the CDSOM used daily mean temperature as opposed to other measures of thermal 493 forcing such as daily maximum and minimum temperature, which some studies have suggested 494 may be better predictors. However, results from CDSOM using daily maximum and minimum 495 496 temperatures as inputs did not show significant differences from results based on daily mean 497 temperatures (not shown), and more generally, results from studies that have explored this question are somewhat inconsistent (c.f., Huang et al., 2020; Piao et al., 2015; Shen et al., 2018). 498 499 That said, because continuous development models are explicitly designed to capture the effects of short-term variability in forcing variables, selection of optimal metrics to this variability is 500 501 clearly important and merits more investigation.

502

503 **5. Conclusions**

504 Changes in springtime phenology are among the most obvious and observable responses of organisms to climate change, but the mechanisms behind these changes are poorly understood 505 506 (Parmesan and Yohe, 2003; Piao et al., 2019). By directly estimating and mapping the 507 geographic dependence of greenup on photoperiod and thermal forcing, results from this study elucidate how the nature and magnitude bioclimatic control on spring phenology depend on 508 geography and climate, and provide a novel and nuanced explanation for why the temperature 509 510 sensitivity of deciduous forests appears to be decreasing. Specifically, our results indicate that apparent changes in temperature sensitivity may reflect a misinterpretation of the data, and where 511 present, observed decreases actually reflect increased dependence on photoperiod. The results 512

513 also help to clarify the mechanisms behind observed changes and have important implications 514 for a variety of ecological processes, such as the role of safety mechanisms that are widely 515 ascribed to photoperiod constraints on spring phenology (Körner and Basler, 2010). For example, Fig. 5 shows that the relative importance of photoperiod decreases as mean annual temperature 516 decreases, which suggests that safety mechanisms related to photoperiod provide only modest 517 518 protection in colder climates (Richardson et al., 2018a). More generally, our results support the 519 argument posited by Zohner et al. (2016) who reported that tree species with strong photoperiod 520 control on leaf-out tend to be located in warmer regions, and challenge the idea that photoperiod 521 provides a safeguard against early leaf emergence in temperate woody species.

522

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527

528 Author contributions

M.M. and M.A.F designed the analysis and led the drafting of the manuscript. M.M. and B.S.
developed the model. MM. performed the analysis. B.S. and A.D.R. contributed analysis ideas
and participated in drafting the manuscript.

532

533 Competing interests

534 The authors declare no competing interests.

535 Appendix

536 **Table A1.** Models descriptions

Model	Model type	Variables	Main characteristics & Statistical assumptions of the model	
CDSOM	Date-driven	$h; \delta h; h_{max}; X; T; L;$ CU; β; P; κ; λ; Y; T _b	Phenological development responses continuously to variations in environmental controls at daily time step throughout pre-season period; Invoking no assumptions about functional relationships between control variables	
TT	Knowledge-driven	$F^*; S_f; T; T_b$	Greenup onset occurs when accumulated forcing reaches a critical threshold, which sorely relies only on thermal forcing with no additional factors	
PTT	Knowledge-driven	$F^*; S_f; T; T_b; L$	Greenup onset occurs when accumulated forcing reaches a critical threshold, but the rate of thermal forcing is regulated by photoperiod	
M1	Knowledge-driven	$F^*; S_f; T; T_b; L; k$	Greenup onset occurs when accumulated forcing reaches a critical threshold, but the rate of thermal forcing is regulated by photoperiod as an exponential	
AT	Knowledge-driven	<i>F</i> *; NCD; <i>a</i> ; <i>b</i> ; <i>c</i>	Greenup onset occurs when accumulated forcing reaches a critical threshold, but the rate of thermal forcing is regulated by the number of chilling days	
CDSOM: Continuous Development Spring Onset Model: TT: Thermal Time model (TT): PTT: Photo-Thermal Time model (PTT):				

537 CDSOM: Continuous Developr nent Spring Onset Model; I: Thermal Time model (11); : Photo-I nal Time model (F M1: Exponential Photo-Thermal Time model (M1); AT: Alternating model; h: latent state; δh : daily latent state increment; h_{max} : 538 theoretical final state of h; X: matrix of predictor variables T, L, and CU (daily mean temperature, day-length, and chilling units, 539 respectively); β : vector of estimated model coefficient for T, L, and CU; P: probability that greenup onset occurs; κ and λ : intercept 540 and slope for logit transformation, respectively; Y: Bernoulli trial indicating whether or not greenup onset has occurred; T_b : base 541 temperature for chilling requirement; F^* : critical threshold that spring greenup onset occurs when the state of forcing (S_f) reaches it; k: 542 exponential coefficient for M1; NCD: number of chilling days; estimated constants for AT 543



546 Fig. A1. Map of the study area. Extents of the US EPA Northern Forest and Eastern Temperate Forest ecoregions, along with the proportion 500 m MODIS pixels labeled as deciduous forests 547 in each grid cell according to the Collection 6 MODIS Land Cover Type product. Red crosses 548 show the USA-National Phenology Network site locations where lilac data are collected. Note 549 that because the MODIS Land Cover Type product uses a threshold of 60% cover to define 550 forest classes, the map shown in Fig. A1 modestly over-represents the actual proportion of 551 552 deciduous forest cover.



Fig. A2. Relationship between anomalies in MODIS onset dates and anomalies in modelpredicted onset dates. Panels (a)-(e) show results for the Continuous Development Spring Onset Model (CDSOM), the thermal time model (TT), the photo-thermal time model (PTT), the exponential photo-thermal time model (M1), and the alternating model (AT), respectively. Dashed lines and correlation coefficients (*r*) show the results from standard major axis regression.





Fig. A3. Relationship between MODIS greenup dates and leaf unfolding dates from the USA-NPN cloned lilac dataset. The colder (red dots) and warmer (blue dots) sites are divided based on mean annual temperature (i.e., colder ≤ 10 °C; warmer > 10 °C). *n* (= 198) is different from the total number of USA-NPN leaf unfolding dates (n = 254) due to cases where no MODIS dates were available because the lilac site was not located in a location dominated by deciduous or mixed forest at the scale of MODIS pixels.



Fig. A4. Relationship between pre-season period length and dependence on thermal forcing.

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776 List of Figure Captions

777 Fig. 1. Model results for a randomly selected grid cell. (a) Relationship between the greenup onset dates from MODIS and onset dates estimated by the model. (b) The distribution of model 778 coefficients for each control variable (i.e., the relative dependence on each climate control; 779 Therm.: thermal forcing; Photo.: photoperiod; Chill.: chilling units). (c) Time series of the latent 780 state (red line) and the length of the pre-season (identified by the horizontal arrow). In panel (a), 781 782 each dot (total n = 100) represents an individual pixel-year sampled from the grid cell comprised 783 of 10 by 10 MODIS pixels across 17 years of the study period (i.e., 100 out of the total 1,700 pixel-years). 784

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Fig. 2. Continuous Development Spring Onset Model (CDSOM) performance. (a) Geographic variation in model root-mean-square errors (RMSE) between greenup onset dates observed from MODIS and onset dates predicted by the CDSOM model. (b) Boxplots showing the distribution of RMSEs for the CDSOM model and four widely used conventional spring greenup models. M1: The exponential photo-thermal time model; PTT: The photo-thermal time model; TT: The thermal time model; AT: The alternating model. In panel (b), boxplots are presented in increasing order of magnitude with respect to mean RMSE.

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Fig. 3. RMSE results across models for anomalous years. (a) Boxplots of RMSEs for each model for 2010 and 2012. (b) Boxplots showing increase in RMSEs for model predictions for all years versus anomalous years (i.e., RMSEs for 2010 and 2012 – RMSEs for 2001-2017) at each grid cell. CDSOM: continuous development spring onset model; M1: The exponential photo-thermal time model; PTT: The photo-thermal time model; TT: The thermal time model; AT: The
alternating model. Boxplots are presented in increasing order of magnitude with respect to mean
RMSE.

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Fig. 4. Geographic variation in the dependence of spring greenup onset date to: (a) thermal forcing, (b) photoperiod, and (c) chilling units. In panel (d), boxplots show the distribution of model coefficients for each control variable during the pre-season period prior to leaf emergence in Northern Forests (blue) versus Eastern Temperate Forests (red). Differences between the means in both cases are statistically significant (p < 0.001).

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Fig. 5. Relative importance (*RI*) of thermal forcing versus photoperiod. Circles in red and blue show locations where thermal forcing and photoperiod, respectively, exert stronger control on the timing of spring greenup; purple circles identify locations where the magnitude of thermal forcing and photoperiod are roughly equivalent. The size of each circle is proportional to the magnitude of *RI* in each cell.

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Fig. 6. Dependence of cloned lilac leaf unfolding date on thermal forcing and photoperiod, and relative importance (*RI*). 254 total leaf unfolding dates from cloned lilac were divided into two groups based on mean annual temperature (≤ 10 °C, n = 182; > 10 °C, n = 72). The left panel plots the mean dependence of leaf unfolding on thermal forcing and photoperiod estimated by the CDSOM. The right panel plots the mean *RI* in each group. Positive *RI* indicates stronger control by thermal forcing relative to photoperiod. Vertical lines show ±1 standard deviation.

Fig. 7. Variation in pre-season period and the relationship between greenup dependence on photoperiod and length of pre-season period. (a) Geographic pattern in pre-season period, and (b) log-log relationship between the dependence of greenup on photoperiod and the length of the pre-season period.

Fig. A1. Map of the study area. Extents of the US EPA Northern Forest and Eastern Temperate Forest ecoregions, along with the proportion 500 m MODIS pixels labeled as deciduous forests in each grid cell according to the Collection 6 MODIS Land Cover Type product. Red crosses show the USA-National Phenology Network site locations where lilac data are collected. Note that because the MODIS Land Cover Type product uses a threshold of 60% cover to define forest classes, the map shown in Fig. A1 modestly over-represents the actual proportion of deciduous forest cover.

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Fig. A2. Relationship between anomalies in MODIS onset dates and anomalies in modelpredicted onset dates. Panels (a)-(e) show results for the Continuous Development Spring Onset Model (CDSOM), the thermal time model (TT), the photo-thermal time model (PTT), the exponential photo-thermal time model (M1), and the alternating model (AT), respectively. Dashed lines and correlation coefficients (*r*) show the results from standard major axis regression.

Fig. A3. Relationship between MODIS greenup dates and leaf unfolding dates from the USA-NPN cloned lilac dataset. The colder (red dots) and warmer (blue dots) sites are divided based on mean annual temperature (i.e., colder ≤ 10 °C; warmer > 10 °C). *n* (= 198) is different from the total number of USA-NPN leaf unfolding dates (n = 254) due to cases where no MODIS dates were available because the lilac site was not located in a location dominated by deciduous or mixed forest at the scale of MODIS pixels.

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847 Fig. A4. Relationship between pre-season period length and dependence on thermal forcing.