

Modeling the effect of temperature on rhizome sprouting in the invasive weed silverleaf nightshade (*Solanum elaeagnifolium*)

Research Article

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Abstract

Silverleaf nightshade (*Solanum elaeagnifolium* Cav.), a noxious, highly invasive perennial weed, poses a significant threat to irrigated summer crops, vegetables, and orchards. This weed has the ability to reproduce both sexually through seed production and asexually via an extensive underground rhizome network, the latter playing a major role in the weed's invasion, establishment, and persistence. Our aims were thus to assess the impact of temperature on rhizome sprouting for fragments of different lengths and to model the sprouting dynamics. The influence of temperature on the sprouting of rhizome fragments (2.5-, 5-, 7.5-, or 10-cm long) was investigated in growth chambers at eight temperatures ranging from 10 to 45 C. The highest sprouting proportions for 10-cm rhizome fragments were recorded at 30 and 35 C in complete darkness. The highest sprouting time for all fragment lengths was observed at 15 C in complete darkness. Modeling sprouting rates as a function of temperature gave the cardinal temperatures for the four different rhizome fragment lengths, with T_b (base temperature) values of 12.80, 9.34, 9.14, and 9.50 C, T_o (optimal temperature) values of 38.90, 36.60, 35.16, and 34.86 C, and T_c (ceiling temperature) values of 39.80, 40.08, 40.50, and 40.80 C for rhizome lengths of 2.5, 5, 7.5, and 10 cm, respectively. Based on these findings, the potential for *S. elaeagnifolium* to spread to new areas and possible new management strategies are discussed; these offer a novel approach for informed decision making regarding the control of this weed.

Introduction

Silverleaf nightshade (*Solanum elaeagnifolium*, Cav.) is a highly invasive weed that has spread rapidly worldwide, infesting both irrigated and rainfed agricultural systems (Karmezi et al. 2022; Krigas et al. 2023). Native to the southwestern United States and northern Mexico, *S. elaeagnifolium* has been introduced into numerous countries, including Greece, Morocco, India, Tunisia, Egypt, and Australia, primarily due to human activities (Roberts and Florentine 2022; Sayari et al. 2022; Uludag et al. 2016). It is a deep-rooted perennial plant that reproduces sexually through seed production and asexually via rhizomes, enabling rapid colonization (Tataridas et al. 2022a). The weed is thus very adaptable and thrives in diverse environments, such as natural reserves, riverbanks, and cultivated fields (Chavana et al. 2021; Mekki 2007). Introduced into Israel in the 1950s, *S. elaeagnifolium* has become a significant weed pest, especially in irrigated summer crops such as cotton (*Gossypium hirsutum* L.), watermelon [*Citrullus lanatus* (Thunb.) Matsum. & Nikai], sorghum [*Sorghum bicolor* (L.) Moench], and maize (*Zea mays* L.), as well as in orchards.

The management of *S. elaeagnifolium* in Israel is a complex and challenging task, mainly due to the limitations of current herbicides: the registered herbicides that are selective and are recommended for *S. elaeagnifolium*, such as aminopyralid and imazapyr, are not suitable for agricultural crops (Wu et al. 2016), and other herbicides that are efficacious, such as glyphosate and glufosinate ammonium, lack selectivity and are therefore suitable for use only in limited control scenarios, such as preplanting preparation (Gitsopoulos et al. 2017). Furthermore, the use of mechanical control methods, including cultivation and mowing, for managing *S. elaeagnifolium* is problematic due to the extensive root network of this weed: the soil disturbance caused by mechanical control methods can result in the breakage of the weed rhizomes, thereby promoting an increase in the spread and density of propagules, both of which

facilitate colonization of new areas (Stanton et al. 2011; Tataridas et al. 2022b; Wu et al. 2016). To effectively manage this weed, it is necessary to elaborate and adopt new methods that require expanded knowledge of its biology and ecology (Chauhan 2022; Westwood et al. 2018); such methods can then be integrated into management strategies based on optimal herbicide and mechanical control applications (Grundy et al. 2003; Krueger-Mangold et al. 2006).

The rhizomes of *S. elaeagnifolium* are the main dispersal organ of this perennial species. However, the literature on the biological and environmental factors that impact the sprouting, establishment, and development of *S. elaeagnifolium* rhizomes is limited, with the available literature focused mainly on the impact of rhizome fragment length on plant establishment. Among those studies is a greenhouse study that assessed shoot, dry mass, and fruit production of *S. elaeagnifolium* plants that were developed from rhizomes with various fragment lengths. This study found that 20-cm fragments gave rise to taller plants compared with those originating from 5-cm and 10-cm fragments (Boyd and Murray 1982). However, this study did not evaluate the impact of fragment length on the sprouting pattern of the weed or the interaction of this biological parameter with environmental factors.

Among the various environmental factors influencing the sprouting dynamics and early establishment of weeds, temperature is one of the most prominent (Cezar Moraes de Aguiar et al. 2022; Peters et al. 2014). Consequently, several studies have concentrated on creating prediction models based on thermal time (TT), specifically growing degree days (GDD). The importance of these models, which quantify accumulated heat units above a minimal threshold, lies in their ability to forecast crucial developmental stages, such as rhizome sprouting, for particular species (Dorado et al. 2009; Loddo et al. 2012). Thus, these thermal models enable a more precise understanding of the effects of temperature on plant growth, development, and sprouting dynamics, thereby facilitating the formulation of targeted management strategies to effectively mitigate the spread and impact of these species. However, the construction of a predictive model necessitates preliminary studies to obtain species-specific biological parameters, such as base (T_b), optimal (T_o), and ceiling (T_c) temperatures, also referred to as cardinal temperature thresholds (Holt and Orcutt 1996; Masin et al. 2010). Based on experimental results, threshold models have been employed to predict rhizome sprouting and seed germination responses to temperature (Bradford and Bello 2022; Holt and Orcutt 1996). Such models are being increasingly utilized to enhance weed management and crop protection strategies. Moreover, their application in agronomic contexts can contribute to a better understanding of invasion ecology in natural ecosystems. These models make it possible to forecast the emergence of weeds under field conditions in various environmental scenarios. This predictive capability facilitates the evaluation and refinement of timing strategies for optimal weed management.

Recently, Kapiluto et al. (2022) developed a comprehensive temperature-based predictive model for *S. elaeagnifolium* seed germination under a wide range of temperatures. The study revealed the cardinal temperatures for the seed germination of this species, but it was limited in that it did not address the impact of temperature on rhizome sprouting characteristics and dynamics, and hence it does not provide any insights about the cardinal temperatures for this dispersal organ. An earlier study evaluated the impact of emergence timing on the growth and development of *S. elaeagnifolium* plants that were generated from rhizomes versus

seeds (Zhu et al. 2013). That study used the cardinal temperatures of $T_b = 10$ C, $T_o = 30$ C, and $T_c = 40$ C for the GDD calculations, but those cardinal temperatures were assumed and not estimated on the basis of experimental work. An additional drawback was that the same values were used for rhizomes and seeds, without taking into consideration differences in germination and sprouting patterns of these two dispersal organs. Therefore, the objectives of the current study on *S. elaeagnifolium* were to: (1) determine the influence of constant temperatures and fragment length on rhizome sprouting, (2) determine the cardinal temperatures for sprouting, and (3) develop prediction models for the sprouting dynamics at different temperatures.

Materials and Methods

Two experiments were conducted, one in May 2020 (Run 1) and the other in August 2020 (Run 2), at the Ne've Ya'ar Research Center, Israel (32.7078°N, 35.1784°E). These experiments aimed to examine the responses of rhizome sprouting to temperature and fragment length and to model the time to rhizome sprouting. The experiments were conducted under temperature- and light-controlled conditions in a growth chamber.

Plant Material and Preliminary Treatments

Solanum elaeagnifolium rhizomes were collected from fields located next to Nahalal in the Jezre'el Valley of Israel (32.42°N, 35.12°E; altitude: 85 m) in May and August of 2020. *Solanum elaeagnifolium* plants >20 cm in height were randomly selected, and their rhizomes were removed from 25 to 35 cm below the soil surface. The rhizomes were washed under tap water, and additive roots and soil debris were removed. Thereafter, the rhizomes were wrapped in a moist cloth to prevent them from drying out and stored under ambient conditions (22 to 25 C). Finally, the rhizomes were cut to fragments 2.5, 5, 7.5, and 10 cm in length, and the plant material was always prepared as described 1 d before use.

Experimental Setups

Rhizome sprouting under different temperature regimes was evaluated by placing the *S. elaeagnifolium* rhizome fragments in pots held in plastic trays (Tivan-Biotec, Kfar-Saba, Israel) as follows. Each tray (54.7 cm long, 27.8 cm wide, and 6.2 cm high) held 12 individual pots (each 12.3 cm long, 7.9 cm wide, and 5.7 cm high) filled with commercial potting medium (Tuff, Marom Golan, Israel). Single fragments, all of the same length (2.5, 5, 7.5, or 10 cm), were placed in each pot of a tray and were then covered with a 1-cm layer of the potting medium. In total, we used 96 trays containing 1,152 rhizome fragments (4 fragment lengths \times 12 pots per tray \times 3 replications \times 8 temperature regimes) for each run. The trays were placed in growth chambers (Conviron Ltd, USA) for 30 d at constant temperatures of 10, 15, 20, 25, 30, 35, 40, or 45 C in complete darkness (0/24-h light/dark). Fifty milliliters of water was added daily for each pot to maintain moisture. Sprouting rhizomes in each pot were counted daily. Rhizomes were considered to have sprouted when the stem was 5 mm or longer and could be observed next to the potting medium surface. Each of the two runs was arranged in a complete randomized design with three replications (trays) for each rhizome length. All statistical analyses were performed using RStudio software in the R environment (R Core Team 2021) and GGLOT2 (Wickham 2016) was used to generate all figures.

Impact of constant temperatures on *Solanum elaeagnifolium* sprouting

The data collected from each tray (containing *S. elaeagnifolium* fragments of a particular length) at each run were used to parameterize a time-to-event model by using the DRCTE package (Onofri et al. 2022). The number of sprouted fragments at the different temperatures was predicted using a three-parameter log-logistic equation:

$$f(t) = \frac{d}{1 + \exp\{b(\log(t) - \log(e))\}} \quad [1]$$

where d is maximum sprouting, e is the time (days) at which 50% of the rhizomes had sprouted, and b is an absolute value proportional to the slope of f at time t . The difference between curves of the two runs in each temperature and fragment length combination was compared using the *compCDF* function (i.e., compare time-to-event curves). The P-value was calculated by the permutation approach, and curves were considered significantly different when the P-value was <0.05 . Additionally, the similarity between the parameters of the log-logistic model from the two runs was determined by pairwise comparison using the *compParm* function. Here, parameters were considered significantly different when the P-value was ≤ 0.05 (Onofri et al. 2022; Ritz et al. 2015).

The fitted log-logistic curves were used to derive the final sprouting proportion (FSP) and the time to sprouting for the 30th percentile of each fragment length (T30). Following Bradford (2002), the T30 percentile was used, because in most combinations of fragment length and temperature, the maximal sprouting proportion (d) was <0.3 . Then, generalized linear models (GLMs) with binomial and logit link were fit to examine the effect of run, rhizome fragment length, and temperature and their interactions on FSP by using the *glm* and *ANOVA* functions. For T30, a similar GLM analysis was done, but with a log-normal error and an identity link. For both analyses, the temperature factor contained six levels, as not all the rhizome fragments sprouted at 10 and 45 C. The rhizome fragment length factor contained four levels, and the run factor contained two levels. Residual plots were employed to assess the assumption of homogeneity of variance. For both GLMs, least-squares means were computed using the *EMMEANS* package (Lenth 2023). Compact letter displays of all pairwise comparison were estimated using the *MULTCOMP* package (Hothorn et al. 2008).

Modeling Sprouting Rate as a Function of Temperature

Sprouting rate (SR) values (the inverse of the time needed for sprouting) derived from the fitted model (Equation 1) were used to develop sprouting prediction models. Such models use the cardinal temperatures, T_b , T_o , and T_c , to quantify the impact of temperatures on rhizome sprouting. Due to the variability of SR across the different fragment lengths, SRs were computed for the different percentiles for each fragment length: 10th, 20th, 30th, 40th, and 50th using the *quantile* function in the DRCTE package. The SRs for the unsprouted fractions were also used in this analysis. In accordance with Bradford (2002), the sprouting data were not normalized relative to the maximal, and the absolute values were employed. Further, Bradford noted different values of maximal SRs in the different percentiles. These observations suggest the use of an addition parameter, S_m , which represents this maximal value of SR. The *DRC.beta* function in the *AOMISC* package (Onofri 2020) was used to fit the derived SR values to a beta function nonlinear

regression model (Yin et al. 1995) correlating sprouting rate and temperature for each fragment length as follows:

If $T_b < T < T_c$, then :

$$SR(T, g) = \frac{1}{t_g} = S_{m(g)} \left\{ \left(\frac{T_{i(g)} - T_{b(g)}}{T_{o(g)} - T_{b(g)}} \times \frac{T_{c(g)} - T}{T_{c(g)} - T_{o(g)}} \right)^{\left(\frac{T_{c(g)} - T_{o(g)}}{T_{o(g)} - T_{b(g)}} \right)} \right\}^{a(g)} \quad [2]$$

If $T \leq T_b$ or $T \geq T_c$, then :

$$SR = 0$$

where SR is the sprouting rate calculated for each fragment length for each percentile and run (g), T is the sprouting temperature, a is a shaping parameter, and S_m is the highest SR that occurs at the T_o . This model predicted no sprouting below T_b or above T_c .

Equation 2 estimates five parameters, and because there are five percentiles, the total number parameters estimated for each fragment length was 25. However, according to Mesgaran (2018), the number of parameters in the prediction model can be reduced by using an equation that sets the same T_b , T_o , and T_c values across the different percentiles, as follows:

If $T_b < T < T_c$, then :

$$SR(T, g) = \frac{1}{t_g} = S_{m(g)} \left\{ \left(\frac{T_i - T_b}{T_o - T_b} \times \frac{T_c - T}{T_c - T_o} \right)^{\left(\frac{T_c - T_o}{T_o - T_b} \right)} \right\}^a \quad [3]$$

If $T \leq T_b$ or $T \geq T_c$, then :

$$SR = 0$$

This model has nine parameters and is based on a positive relationship between the percentiles and the value of SR. Two more modeling approaches were also tested based on Equation 3. The first—with 13 parameters (Bradford 2002)—assumes fixed T_b and T_o parameters and a variable T_c parameter across the different percentiles. The second—with 13 parameters—assumes the fixed T_b and T_c parameters and a variable T_o parameter across the percentiles (Mesgaran et al. 2017). For simplicity, we will refer to our four models as follows: the 25-parameter model is designated the “variable model,” the 9-parameter model is designated the “fixed T_c model,” and the two 13-parameter models are designated the “variable T_c model” and the “variable T_o model.” To determine potential data pooling between the two runs, the 95% confidence interval (CI) values of the estimated cardinal temperatures and a (the shape parameter) were compared for all four models.

Finally, the Akaike information criterion (AIC) was used to assess the models. This method compares the goodness of fit of the models; the model with the lowest AIC value was selected. The AIC method incorporates the amount of reduction of the residual sum of squares (RSS) and the model complexity (Burnham and Anderson 2004):

$$AIC = 2n \ln \left(\frac{RSS}{n} \right) + 2k \quad [4]$$

where n is the number of observations, and k is the number of parameters used in the model.

Development of the Sprouting Dynamics Model

To develop a TT sprouting predictive model among fragment lengths at all temperatures, the cardinal temperatures from the favorable model—according to the AIC criterion—were used for computing GDD. The accumulated values over time (n) were estimated for each rhizome length using the following equation (Cochavi et al. 2016; Mesgaran 2018):

$$\text{GDD} = \sum_{i=1}^n \left\{ \left(\frac{T_i - T_b}{T_o - T_b} \times \frac{T_c - T}{T_c - T_o} \right)^a \right\} (T_o - T_b) \quad [5]$$

where T_i is the daily temperature for the i th day, and n is the total number of days for which GDD are to be calculated. Equation 5 takes into consideration the sub- and supra-optimal temperature ranges in the GDD accumulation. The response of *S. elaeagnifolium* sprouting to GDD was analyzed using a four-parameter Weibull equation:

$$\text{cumulative sprouting} = a \times \left\{ 1 - \exp \left[- \left(\frac{\text{GDD} - \text{Lag}}{b} \right)^c \right] \right\} \quad [6]$$

where a is the maximal sprouting, c is the shape parameter that determines the skewness and kurtosis of the equation, b is the scale parameter regardless of the shape value, and lag is the estimate of the time required for the sprouting of the first rhizome. The root-mean-square error (RMSE) was calculated as an indicator of goodness of fit (Lati et al. 2011; Mobli et al. 2022).

Results and Discussion

Impact of Constant Temperatures on *Solanum elaeagnifolium* Sprouting

The sprouting of *S. elaeagnifolium* was adequately described using a log-logistic time-to-event model for all temperature regimes for each fragment length (Figure 1). However, the curve compression revealed significant differences between the two runs (P -value < 0.05) in 11 out of the 24 combinations of temperatures and fragment length model (Figure 1). Furthermore, the pairwise compression of the parameters derived from these 11 models also revealed significant differences (Supplementary Table S1). Thus, at this stage of the model development, the data were not pooled, and each run was analyzed separately.

In both runs, there was no sprouting at the two extreme temperature regimes (10 and 45 C), but sprouting was observed starting from a temperature of 15 C, and the maximum sprouting (parameter d) was found to increase with fragment length. For example, in Run 1, this parameter reached values of 0.19, 0.35, 0.47, and 0.67 for the 2.5-, 5-, 7.5-, and 10-cm lengths, respectively, at 15 C (Table 1). Also in Run 1, the highest sprouting proportion was observed at 35 C, with the maximum sprouting reaching values of 0.79, 0.83, 0.90, and 1.00 for the 2.5-, 5-, 7.5-, and 10-cm lengths, respectively. In both runs, a decline in maximum sprouting was observed at 40 C, and for the 10-cm fragment length in Run 2, the value of this parameter dropped to 0.64 (Table 1). The time taken for 50% of the rhizomes to sprout (e parameter) was found to decrease with length for shorter versus longer fragments. In Run 2, for example, this parameter was the highest for all fragments

at 15 C, reaching values of 23.8, 16.8, 17.1, and 16.1 d for the 2.5-, 5-, 7.5-, and 10-cm fragment lengths, respectively (Table 1). The e parameter declined as the temperature increased, with the lowest values being obtained at 30 and 35 C. In Run 1, the value of this parameter was 7.8 and 5.4 d for fragment lengths of 2.5 and 5 cm, respectively, while for the longer fragments of 7.5 and 10 cm, it was 4.1 and 3.8 d, respectively. The GLM analysis of the T30 values also revealed that the sprouting times of the longer fragments were significantly lower than those for the shorter ones in both runs (Table 2). For example, in Run 2, the T30 values at 25 C for the 7.5- and 10-cm fragments were ~7 and ~6 d, respectively. These values were significantly lower than those for the shorter fragments of 2.5 and 5 cm, with T30 values of 10.3 and 15.5 d, respectively. The shortest time to sprouting in Run 1 (i.e., 4.5 d) was observed at 35 C for the fragment length of 7.5 cm, while the shortest time to sprouting in Run 2 (i.e., 4.3 d) was observed at 30 C for fragments of 10 cm (Table 2).

Our results show the wide range of temperatures, namely, 15 to 40 C, under which *S. elaeagnifolium* rhizomes can sprout, with optimal sprouting occurring at the relatively high temperatures of 30 and 35 C. Similar results have been reported for other perennial weed species with subsoil dispersal organs that are typical for Mediterranean climate regions, such as bermudagrass [*Cynodon dactylon* (L.) Pers.] at 33 C (Satorre et al. 1996) and yellow nutsedge (*Cyperus esculentus* L.) at 28 C (Li et al. 2000), indicating the preference for high temperatures for sprouting of such plants. Furthermore, our results indicate that the maximum sprouting of *S. elaeagnifolium* rhizomes was higher by 10 C than that for seed germination (Kapiluto et al. 2022). In general, rhizomes contain larger carbohydrate reserves than seeds, which in turn enable them to survive and sprout in extreme environments (Anbari et al. 2011; Chen et al. 2015; Mangoale and Afolayan 2020; Yu et al. 2001). The rhizomes thus provide *S. elaeagnifolium*, like other perennial species, with a substantial advantage in terms of adaptability and colonization potential under Mediterranean and extreme climates (Travlos 2013; Uludag et al. 2016). In addition, we also observed that the longer rhizome fragments sprouted more rapidly, with higher FSP values (Table 2), regardless of temperature. These results can also be attributed to the larger carbohydrate reserves in the longer fragments. It is also likely that the longer rhizomes contain more buds, which contribute to the higher FSP values.

Modeling Sprouting Rate as a Function of Temperature

The fitted curves were used to derive the sprouting rates for the 10th, 20th, 30th, 40th, and 50th percentiles, which were used to parameterize Equation 1. By regarding the percentile g as a factor of the rhizome fragment (fragment size), we reached a total number of 25 parameters. In agreement with previous germination modeling studies, we tested three other modeling approaches with 9 (Washitani 1987) and 13 (Bradford 2002; Mesgaran et al. 2017; Rowse and Finch-Savage 2003) parameters that enable the constant cardinal temperatures to be varied for all percentiles. Here, the 95% CI analysis revealed no significant differences between the estimates of the three cardinal temperatures and the shape parameters of the two runs. This trend was revealed in all four tested models, and thus, at this stage of the model development, data from the two runs were pooled (Supplementary Figure S1).

The computed AIC showed some differences between the accuracy levels of the four models. As Table 3 shows, the nine-parameter fixed T_c model provided the best fit, with the smallest AIC value in all rhizome fragment lengths, with the extracted

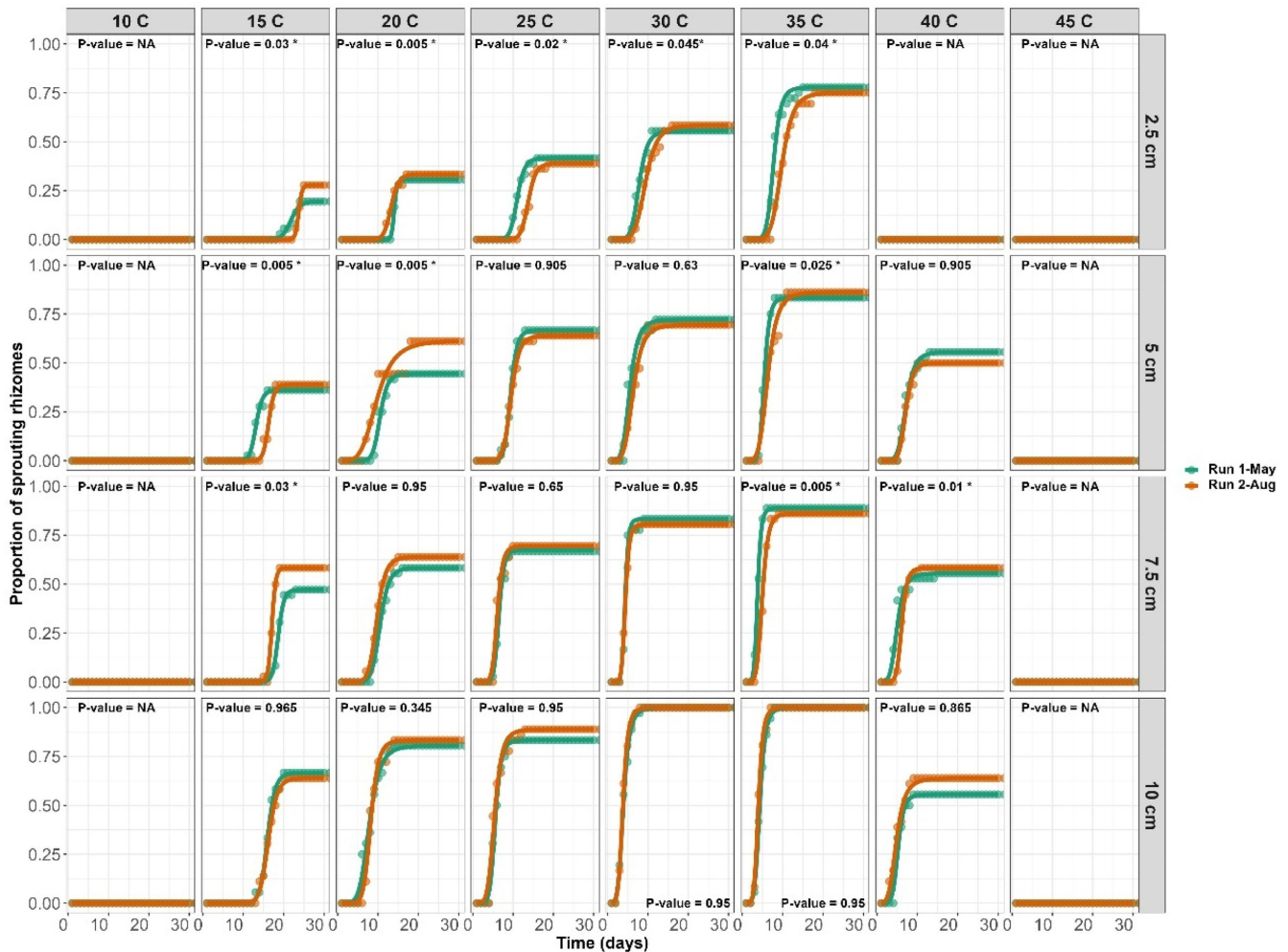


Figure 1. Time-to-event, three-parameter log-logistic equation: $f(t) = d / (1 + \exp\{b(\log(t) - \log(e))\})$ showing the relationship between constant temperature (C) and sprouting proportion of *Solanum elaeagnifolium* rhizomes of four different fragment lengths (2.5, 5, 7.5, and 10 cm) in the two runs, where d is the maximum sprouting, e is the time (days) at which 50% of the rhizomes had sprouted, and b is an absolute value proportional to the slope of f at time t . Coefficients for the equation parameters are presented in Table 1; $n = 3$. The P-value was calculated by the permutation approach to determine the difference between curves of the two runs. Curves were considered significantly different when the P-value was < 0.05 .

values being: $-1,309$, $-1,215$, $-1,191$, and $-1,219$ for the rhizome lengths of 2.5, 5, 7.5, and 10 cm, respectively (Table 3). Thus, the fixed T_c model was used to determine the cardinal temperatures of the pooled data of the rhizome fragment lengths. The T_b values were 12.80 ± 1.01 , 9.34 ± 0.71 , 9.14 ± 1.87 , and 9.50 ± 0.68 , the T_o values were 38.90 ± 0.07 , 36.60 ± 0.23 , 35.16 ± 0.57 , and 34.86 ± 0.40 , and the T_c values were 39.80 ± 0.07 , 40.08 ± 0.03 , 40.50 ± 0.40 and 40.80 ± 0.40 for the rhizome lengths of 2.5, 5, 7.5, and 10 cm, respectively. In addition, the S_m parameter decreased with the percentiles in all fragment lengths (Table 4). The values of the S_m parameter increased as the fragment length increased: for the 2.5-cm fragment length, this value was 0.18, 0.16, 0.14, 0.11, and 0.08, while for the 10-cm fragment length, this value was 0.36, 0.32, 0.29, 0.26, and 0.24 for the 10th, 20th, 30th, 40th, and 50th percentiles, respectively.

The modeling approaches used in this study take into consideration the different sprouting characteristics of the various percentiles within the rhizome fragment lengths (Figure 2). These differences were reflected by the S_m parameter in Equation 3, which represents the maximal value of SR. Understanding the sprouting dynamics among different percentiles of *S. elaeagnifolium* allows

for informed decision making in control activities. This insight enables precise targeting of control measures to the timing when the majority of the fragment has emerged, preventing both early and late applications.

The beta-function model was then used to predict *S. elaeagnifolium* sprouting rate among all rhizome lengths and was shown to be suitable for predicting the effects of temperature on the sprouting rate of this weed. The model's effectiveness in predicting cardinal temperatures and the development of phenological events in various plant species has indeed been demonstrated previously (Cochavi et al. 2016, 2018; Yin et al. 1995). Kapiluto et al. (2022), who performed a study similar to the current one, but for seeds, revealed cardinal temperatures of 10.8, 23.8, and 35.9 C for T_b , T_o , and T_c , respectively. It may be seen that these values for *S. elaeagnifolium* seeds and rhizomes differed mainly for T_o and T_c , which were higher by 12 and 5 C, respectively, among the rhizome fragments. These results further emphasize the plasticity of *S. elaeagnifolium* and its ability to adapt to temperature variations, similar to other invasive species, and the importance of conducting studies focusing on rhizomes (Clements and DiTommaso 2011; Peters et al. 2014). It can be assumed that

Table 1. Coefficients and their standard errors (in parentheses) and P-values describing the relationship between sprouting at different constant temperature (C) regimes^a and rhizome fragment lengths of *Solanum elaeagnifolium* in the two runs

Length cm	Temp C	Run	Coefficient					
			<i>b</i>	P-value (<i>b</i>)	<i>d</i>	P-value (<i>d</i>)	<i>e</i>	P-value (<i>e</i>)
2.5	15	1	-16.66 (6.62)	0.012	0.19 (0.06)	0.003	22.91 (0.21)	<0.001
	15	2	-56.94 (19.34)	0.003	0.27 (0.07)	0.003	23.87 (0.18)	<0.001
	20	1	-31.28 (14.12)	0.027	0.30 (0.07)	<0.001	13.86 (0.14)	<0.001
	20	2	-13.91 (3.57)	<0.001	0.32 (0.076)	<0.001	13.02 (0.37)	<0.001
	25	1	-12.62 (2.94)	<0.001	0.41 (0.08)	<0.001	11.16 (0.38)	<0.001
	25	2	-13.55 (3.28)	<0.001	0.39 (0.080)	<0.001	13.88 (0.38)	<0.001
	30	1	-7.61 (1.46)	<0.001	0.55 (0.08)	<0.001	7.82 (0.41)	<0.001
	30	2	-7.75 (1.5)	<0.001	0.58 (0.081)	<0.001	8.83 (0.41)	<0.001
	35	1	-7.98 (1.4)	<0.001	0.79 (0.06)	<0.001	7.61 (0.24)	<0.001
	35	2	-6.82 (1.18)	<0.001	0.74 (0.072)	<0.001	9.43 (0.38)	<0.001
40	1	—	—	—	—	—	—	—
	2	—	—	—	—	—	—	—
5	15	1	-15.66 (4.05)	0.001	0.35 (0.07)	<0.001	13.00 (0.21)	<0.001
	15	2	-22.64 (5.53)	0.001	0.37 (0.08)	<0.001	16.89 (0.18)	<0.001
	20	1	-11.27 (2.39)	<0.001	0.44 (0.08)	<0.001	10.45 (0.39)	<0.001
	20	2	-4.52 (0.92)	<0.001	0.62 (0.08)	<0.001	9.62 (0.82)	<0.001
	25	1	-12.21 (2.17)	<0.001	0.67 (0.07)	<0.001	9.42 (0.28)	<0.001
	25	2	-9.37 (1.72)	<0.001	0.65 (0.08)	<0.001	9.38 (0.32)	<0.001
	30	1	-5.95 (1.04)	<0.001	0.74 (0.07)	<0.001	5.41 (0.33)	<0.001
	30	2	-5.43 (0.97)	<0.001	0.7 (0.08)	<0.001	6.26 (0.42)	<0.001
	35	1	-8.44 (1.37)	<0.001	0.83 (0.06)	<0.001	5.43 (0.23)	<0.001
	35	2	-5.29 (0.86)	<0.001	0.87 (0.06)	<0.001	6.21 (0.35)	<0.001
40	1	-6.82 (1.33)	<0.001	0.56 (0.03)	<0.001	6.98 (0.35)	<0.001	
	2	-9.23 (1.91)	<0.001	0.50 (0.08)	<0.001	6.8 (0.29)	<0.001	
7.5	15	1	-20.28 (5.52)	0.002	0.47 (0.08)	<0.001	18.86 (0.15)	<0.001
	15	2	-34.31 (7.26)	0.002	0.58 (0.08)	<0.001	17.18 (0.2)	<0.001
	20	1	-11.70 (2.34)	<0.001	0.58 (0.08)	<0.001	10.30 (0.30)	<0.001
	20	2	-15.21 (2.75)	<0.001	0.64 (0.08)	<0.001	9.64 (0.25)	<0.001
	25	1	-10.04 (1.83)	<0.001	0.68 (0.07)	<0.001	6.52 (0.22)	<0.001
	25	2	-9.05 (1.67)	<0.001	0.7 (0.08)	<0.001	6.00 (0.24)	<0.001
	30	1	-10.12 (1.89)	<0.001	0.83 (0.06)	<0.001	4.29 (0.16)	<0.001
	30	2	-9.43 (1.63)	<0.001	0.81 (0.07)	<0.001	4.45 (0.19)	<0.001
	35	1	-8.68 (1.47)	<0.001	0.90 (0.05)	<0.001	3.79 (0.17)	<0.001
	35	2	-6.94 (1.15)	<0.001	0.87 (0.06)	<0.001	4.89 (0.25)	<0.001
40	1	-4.84 (1.11)	<0.001	0.55 (0.08)	<0.001	5.04 (0.31)	<0.001	
	2	-8.99 (1.78)	<0.001	0.59 (0.08)	<0.001	6.09 (0.25)	<0.001	
10	15	1	-18.56 (3.2)	<0.001	0.67 (0.07)	<0.001	16.04 (0.32)	<0.001
	15	2	-16.57 (2.88)	<0.001	0.64 (0.08)	<0.001	16.13 (0.37)	<0.001
	20	1	-5.70 (0.95)	<0.001	0.81 (0.06)	<0.001	7.68 (0.43)	<0.001
	20	2	-9 (1.46)	<0.001	0.83 (0.062)	<0.001	7.93 (0.25)	<0.001
	25	1	-8.03 (1.30)	<0.001	0.83 (0.06)	<0.001	5.60 (0.25)	<0.001
	25	2	-5.84 (0.95)	<0.001	0.89 (0.05)	<0.001	5.3 (0.25)	<0.001
	30	1	-5.88 (0.99)	<0.001	1.00 (0.0)	<0.001	3.80 (0.18)	<0.001
	30	2	-5.94 (0.97)	<0.001	1.00 (0.0)	<0.001	3.81 (0.20)	<0.001
	35	1	-6.54 (1.09)	<0.001	1.00 (0.0)	<0.001	4.18 (0.18)	<0.001
	35	2	-6.72 (1.11)	<0.001	1.00 (0.0)	<0.001	4.0 (0.17)	<0.001
40	1	-7.07 (1.46)	<0.001	0.55 (0.08)	<0.001	5.07 (0.24)	<0.001	
	2	-6.42 (1.21)	<0.001	0.67 (0.07)	<0.001	4.34 (0.24)	<0.001	

^aA time-to-event, three-parameter log-logistic equation was used. Equation 1: $f(t) = d / \{1 + \exp(b(\log(t) - \log(e)))\}$, where *d* is the maximum sprouting, *e* is the time (days) at which 50% of the rhizomes had sprouted, and *b* is an absolute value proportional to the slope of *f* at time *t*.

the higher carbohydrate content in rhizome buds, compared with seeds, contributes to increased resistance to extreme temperatures. In light of the global warming trend, the implications of these results suggest that this species may colonize new areas that are currently characterized by moderate climates (Gioria and Pyšek 2017). It can be assumed that higher temperatures will affect seed germination in *S. elaeagnifolium* and that dispersal via rhizomes will become the dominant path.

Development of the Sprouting Dynamics Model

Following the extraction of the cardinal temperatures, a TT model for estimating cumulative *S. elaeagnifolium* sprouting was developed, using the pooled data for all temperature regimes

and rhizome fragment lengths. Time (in days) was converted to GDD using Equation 5, and the cardinal temperatures derived from the fixed *T_c* model were applied to the pooled data. As Figure 3 shows, the four-parameter Weibull equation adequately described the sprouting dynamics for *S. elaeagnifolium* with a RMSE of 0.154 and P-value of <0.001 for all derived model parameters (Table 5). These results indicate the accurate sprouting prediction ability offered by the cardinal temperatures that were derived from the fixed *T_c* model.

Based on the GDD computation (Equation 5) and the parameters that were determined via Equation 6, it was predicted that *S. elaeagnifolium* rhizomes have a lag time of 12 GDD, the 50% sprouting would be reached after almost ~30 GDD, and the

Table 2. Influence of temperature (C) and rhizome fragment length (cm) on final sprouting proportion (FSP) and time to 30% sprouting (T30) for *Solanum elaeagnifolium* in the two runs with the 95% confidence intervals given in parentheses^a

Temp C	Length cm	T30 days	
		Run 1	Run 2
15	2.5	NA	25.34 (22.06–28.84) a
20	2.5	15.74 (13.95–17.64) a	16.01 (14.14–17.99) b
25	2.5	13.00 (11.67–14.41) b	15.52 (14.01–17.11) b
30	2.5	8.98 (7.87–10.15) c	10.86 (9.61–12.20) c
35	2.5	8.34 (7.28–9.48) c	10.37 (9.14–11.68) c
40	2.5	NA	NA
15	5	15.15 (13.29–17.13) a	17.56 (15.56–19.68) a
20	5	12.10 (10.74–13.54) b	10.57 (9.30–11.93) b
25	5	10.23 (8.98–11.57) bc	10.37 (9.11–11.71) b
30	5	6.19 (5.24–7.22) d	7.01 (5.99–8.11) c
35	5	6.07 (5.13–7.10) d	6.71 (5.72–7.79) c
40	5	8.37 (7.24–9.57) c	8.38 (7.26–9.59) c
15	7.5	19.93 (18.18–21.76) a	18.09 (16.42–19.83) a
20	7.5	11.52 (10.19–12.92) b	10.85 (9.56–12.22) b
25	7.5	7.40 (6.35–8.54) c	6.96 (5.94–8.06) cd
30	7.5	5.13 (4.28–6.07) d	5.20 (4.34–6.14) e
35	7.5	4.49 (3.70–5.36) d	5.54 (4.65–6.52) de
40	7.5	7.33 (6.29–8.46) c	7.45 (6.40–8.59) c
15	10	16.82 (15.21–18.51) a	17.18 (15.55–18.88) a
20	10	8.18 (7.07–9.38) b	8.87 (7.71–10.11) b
25	10	6.22 (5.27–7.26) cd	5.97 (5.04–6.98) cd
30	10	4.56 (3.77–5.44) e	4.35 (3.58–5.21) e
35	10	4.80 (3.99–5.71) de	4.58 (3.79–5.47) de
40	10	6.43 (5.46–7.49) c	6.98 (5.75–8.34) bc
		FSP	
		Run 1	Run 2
15	2.5	0.20 (0.18–0.21) a	0.28 (0.26–0.30) a
20	2.5	0.31 (0.29–0.33) b	0.33 (0.31–0.35) b
25	2.5	0.42 (0.39–0.44) c	0.39 (0.37–0.41) c
30	2.5	0.56 (0.53–0.58) d	0.58 (0.56–0.61) d
35	2.5	0.78 (0.76–0.80) e	0.75 (0.73–0.77) e
40	2.5	0.00 (0.00–0.00) f	0.00 (0.00–0.00) f
15	5	0.36 (0.34–0.38) a	0.38 (0.36–0.41) a
20	5	0.44 (0.42–0.47) b	0.61 (0.59–0.63) b
25	5	0.67 (0.65–0.69) c	0.64 (0.62–0.67) b
30	5	0.73 (0.71–0.75) d	0.69 (0.67–0.71) c
35	5	0.83 (0.81–0.85) e	0.86 (0.85–0.88) d
40	5	0.56 (0.53–0.58) f	0.50 (0.48–0.52) e
15	7.5	0.47 (0.45–0.50) a	0.58 (0.56–0.61) a
20	7.5	0.58 (0.56–0.61) b	0.64 (0.62–0.66) b
25	7.5	0.66 (0.64–0.69) c	0.69 (0.67–0.71) c
30	7.5	0.83 (0.81–0.85) d	0.80 (0.79–0.82) d
35	7.5	0.89 (0.87–0.90) e	0.86 (0.85–0.88) e
40	7.5	0.56 (0.53–0.58) b	0.58 (0.56–0.61) a
15	10	0.66 (0.64–0.69) a	0.64 (0.62–0.66) a
20	10	0.80 (0.79–0.82) b	0.83 (0.81–0.85) b
25	10	0.83 (0.81–0.85) b	0.89 (0.87–0.90) c
30	10	1.00 (1.00–1.00) c	1.00 (1.00–1.00) d
35	10	1.00 (1.00–1.00) c	1.00 (1.00–1.00) d
40	10	0.56 (0.53–0.58) d	0.64 (0.62–0.66) a

^aValues followed by the same letter are not significantly different according to multiple-comparison procedure with multiplicity adjustment ($P \leq 0.05$).

maximum sprouting would be recorded at 80 GDD (Figure 3). Kapiluto et al. (2022) showed that for seed germination, the GDD values of the lag, 50% germination, and the maximum germination stages were higher, ~50, 140, and 180 GDD, respectively. These results further emphasize that rhizomes, as the main *S. elaeagnifolium* dispersal organ, contribute to the aggressiveness of this weed in agrosystems by rapid sprouting and establishment. The employment of a beta function for computing the GDD to

Table 3. Akaike information criterion (AIC), degrees of freedom (df), and number of parameters of the four models used (references in parentheses) to describe the relationship between sprouting rates and temperature of *Solanum elaeagnifolium* rhizome fragments of four different lengths

	Tested model ^a			
	Fixed T_c (Washitani 1987)	Variable T_c (Bradford 2002)	Variable T_o (Mesgaran et al. 2017)	Variable T_b, T_o, T_c (Parmoon et al. 2015)
Rhizome length				
2.5 cm	-1,309	-1,082	-1,197	-1,185
5 cm	-1,215	-1,181	-1,178	-1,204
7.5 cm	-1,191	-1,175	-1,181	-1,102
10 cm	-1,219	-1,155	-1,216	-1,155
df	10	10	14	26
No. of parameters	9	13	13	25

^a T_c , ceiling temperature, T_o , optimum temperature, T_b , base temperature.

estimate the sprouting rate of *S. elaeagnifolium* as related to TT allows the use of data obtained from the entire examined range of temperatures. To the best of our knowledge, this is the first time a beta function has been used to predict *S. elaeagnifolium* based on TT. The ability to predict rhizome sprouting patterns and dynamics in relation to GDD is an important biological tool that can facilitate optimized decision making for weed control timing; for example, the outcomes of cultivation and herbicide applications can be improved by leveraging their optimal timing estimated by modeling (Grundy 2003). However, to improve its practical application in the field, further research is required. One notable factor to consider is that the rhizomes remain buried in the soil, consistently exposed to local environmental conditions throughout various seasons. There is a gap in determining the initial timing for temperature measurement and GDD accumulation. Eizenberg et al. (2004) approach this gap by proposing January 1 as the starting point for temperature measurement, hypothesizing that this period represents the coldest time of the year, when seeds are still dormant. This date could serve as a reference point in our system, particularly when rhizomes are dormant during low winter temperatures and initiate sprouting as temperatures rise.

The effectiveness of herbicides and mechanical tools, such as cultivation and harrowing, is markedly impacted by the weed phenology stage at the time of application. When weed control tactics are applied under favorable conditions (i.e., young buds), low herbicide rates and mechanical treatments can still provide satisfactory results (Kolberg et al. 2018; Lati et al. 2012). Our findings regarding the phenology dynamics of *S. elaeagnifolium* can also be linked to—and improve—other practical weed control concepts, such as the critical period for weed control (CPWC), that is, the specific stage in a crop’s growth cycle at which controlling weeds is crucial to prevent yield loss (Knezevic et al. 2002). The competition between weed and crop is influenced by temperature conditions, and our findings can improve any CPWC model. Knowledge of the CPWC can enable farmers and land managers to effectively plan and execute weed control measures, ultimately leading to higher crop yields and improved ecosystem health (Knezevic and Datta 2015).

In conclusion, our findings have important implications for predicting the future invasiveness of *S. elaeagnifolium* rhizomes under varying temperature conditions. Understanding the patterns

Table 4. Influence of temperature on the sprouting rates for *Solanum elaeagnifolium* for the different percentiles of four different lengths for the pooled data

Parameter estimate ^a	Rhizome fragment length			
	2.5 cm	5 cm	7.5 cm	10 cm
<i>a</i>	0.9 (0.11)	1.15 (0.09)	1.76 (0.35)	1.37 (0.13)
<i>S_m</i> (10th)	0.18 (0.006)	0.24 (0.005)	0.31 (0.007)	0.36 (0.005)
<i>S_m</i> (20th)	0.16 (0.006)	0.21(0.005)	0.28 (0.006)	0.32 (0.005)
<i>S_m</i> (30th)	0.14 (0.006)	0.19 (0.005)	0.26 (0.006)	0.29 (0.005)
<i>S_m</i> (40th)	0.11 (0.006)	0.17 (0.005)	0.24 (0.006)	0.26 (0.005)
<i>S_m</i> (50th)	0.08 (0.006)	0.14 (0.005)	0.21 (0.006)	0.24 (0.005)
<i>T_b</i> (C)	12.80 (1.01)	9.34 (0.71)	9.14 (1.87)	9.50 (0.68)
<i>T_o</i> (C)	38.90 (0.07)	36.60 (0.23)	35.16 (0.57)	34.86 (0.4)
<i>T_c</i> (C)	39.80 (0.07)	40.08 (0.03)	40.50 (0.4)	40.80 (0.4)

^aParameter estimates for Equation 3: *T_b*, base temperature; *T_o*, optimum temperature; *T_c*, ceiling temperature; and the shape parameter of the equation that gives more flexibility to the model (*a*).

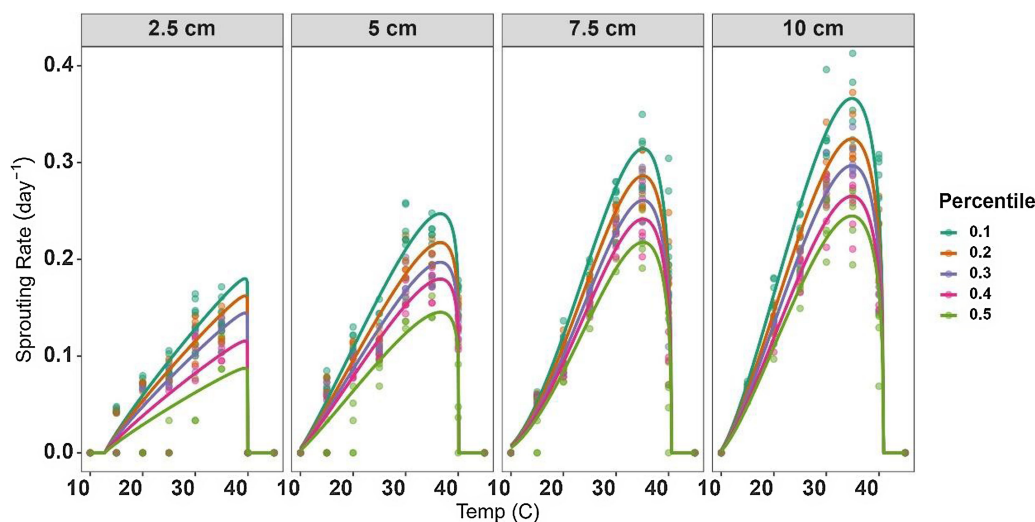


Figure 2. Relationship between sprouting rate (SR) of 10th, 20th, 30th, 40th, and 50th percentiles and temperature for *Solanum elaeagnifolium* rhizomes of four different fragment lengths (2.5, 5, 7.5, and 10 cm). Symbols show the observed data, while lines show the fitted curves (Equation 3) with the parameters given in Table 3.

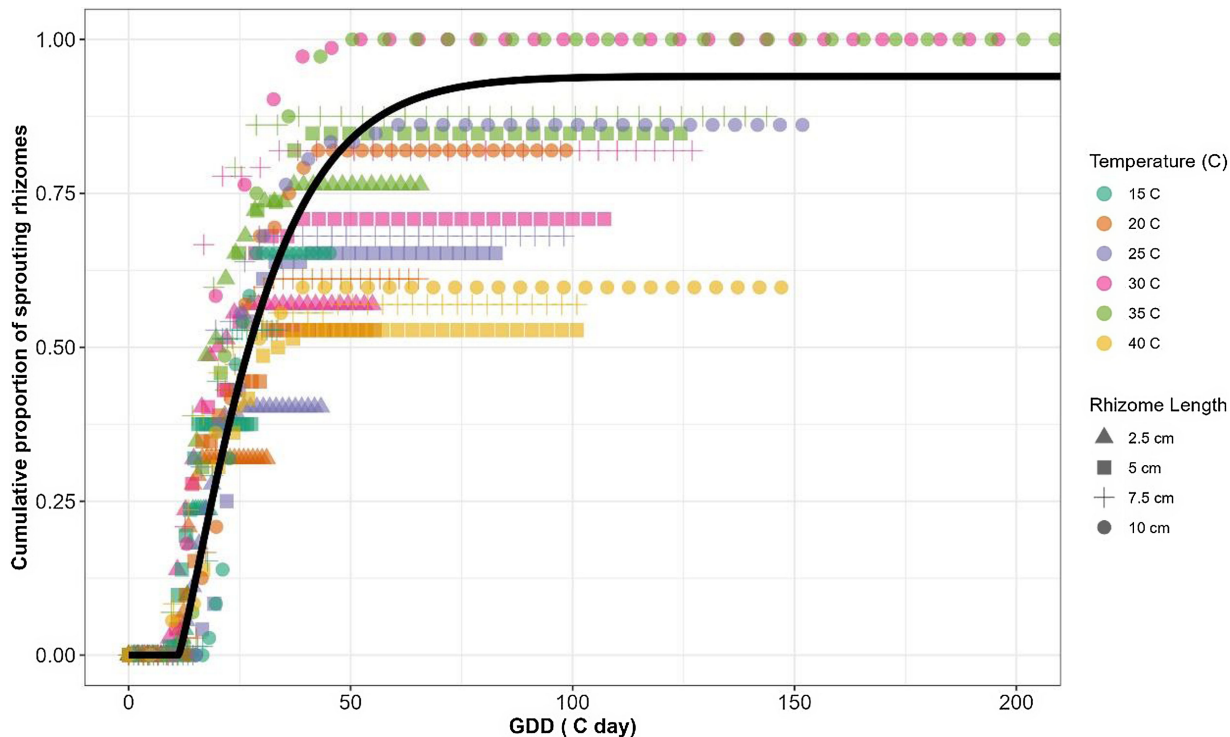


Figure 3. Sprouting dynamics of *Solanum elaeagnifolium* in relation to growing degree days (GDD) obtained using a four-parameter Weibull equation: $f(GDD) = a[1 - \exp(-\{[(GDD - lag)/b]^c\})]$, where *a* is maximal sprouting proportion, *b* is scale parameter regardless of the shape value, *c* is shape parameter that determines the skewness and kurtosis of the equation, and lag is estimate of the time required for the sprouting of the first rhizome. RMSE, root-mean-square error.

Table 5. Four-parameter Weibull equation describing the sprouting dynamics of *Solanum elaeagnifolium* in relation to growing degree days (GDD): cumulative sprouting = $a\{1 - \exp[-(\frac{GDD-lag}{b})^c]\}$, where a is the maximal *Solanum elaeagnifolium* sprouting, b is the scale parameter regardless of the shape value, c is the shape parameter that determines the skewness and kurtosis of the equation, and lag is the estimate of the time required for the sprouting of the first rhizome

Parameter	Estimate	P-value
a	0.92 (0.02)	<0.001
b	30.74 (1.05)	<0.001
c	0.9 (0.05)	<0.001
lag	11.4 (0.4)	<0.001
RMSE ^a	0.154	

^aRMSE, root-mean-square error.

of sprouting in *S. elaeagnifolium* is crucial for optimizing strategies to manage and control this weed. Here, we provide the first comprehensive analysis of its sprouting dynamics by using a precise beta-function model. The cardinal temperatures for sprouting were determined: T_b values were 12.80 ± 1.01 , 9.34 ± 0.71 , 9.14 ± 1.87 , and 9.50 ± 0.68 , the T_o values were 38.90 ± 0.07 , 36.60 ± 0.23 , 35.16 ± 0.57 and 34.86 ± 0.40 , and the T_c values were 39.80 ± 0.07 , 40.08 ± 0.03 , 40.50 ± 0.40 , and 40.80 ± 0.40 for rhizome lengths of 2.5, 5, 7.5, and 10 cm, respectively. Importantly, these cardinal temperature values can be used to predict other phenological stages, thereby enhancing the implementation of weed control measures during later growth stages. As we strive for the adoption of new integrated weed management programs, it is vital to have the comprehensive biological data that will enable optimal outcomes. Further studies conducted under field conditions are thus critical to refine predictive emergence models developed using data obtained under laboratory conditions and to fine-tune weed control timing.

Supplementary material. The supplementary material for this article can be found at <https://doi.org/10.1017/wsc.2024.8>.

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