

# Plant Cell as a Computational Unit: Signal Integration and Decision Making from Root to Flower

Markus Schneider\*

Department of Plant Systems Biology, Universidad Autónoma Chapingo, Texcoco, México

\*Corresponding author: Markus Schneider, [schneider12@chapingo.edu.mx](mailto:schneider12@chapingo.edu.mx)

## Abstract

The traditional view of plants as passive, reactive organisms has been fundamentally overturned. Modern plant science reveals a sophisticated entity capable of sensing, processing, and responding to a multitude of concurrent environmental and internal signals with remarkable precision. This article proposes and elaborates on a paradigm-shifting framework: conceptualizing the plant cell, and by extension, the plant as a whole, as a distributed biological computational system. We argue that plant cells function as fundamental processing nodes within a networked organism, executing complex operations of signal perception, integration, memory, and decision-making that govern development, growth, and survival. This review synthesizes evidence across scales—from molecular networks within single cells to long-distance communication between organs. We detail the cellular “hardware”: receptors as sensors, second messengers as signal transducers, and protein-phosphorylation/ gene-regulatory networks as logic gates. We then explore how specialized tissues, such as the root apex and shoot apical meristem, act as central processing hubs, integrating local and systemic data to direct exploratory growth (root) or developmental fate (shoot). A dedicated figure illustrates this plant-wide “computational network,” while a comprehensive table maps biological components to computational analogs. The article further examines how plants perform cost-benefit analyses and predictive modeling, for instance, in balancing root foraging against shoot growth under resource limitation. Finally, we discuss the implications of this computational perspective for predictive agriculture, such as designing crops with enhanced environmental decision-making logic or using *in silico* models to simulate plant responses. By adopting this lens, we not only deepen our understanding of plant intelligence but also unlock novel bio-inspired computational strategies and engineering approaches for building resilient agricultural systems.

## Keywords

Plant Cell Computation, Signal Integration, Systems Biology, Plant Neurobiology (Analog), Long-Distance Signaling, Root Apex, Shoot Apical Meristem, Phenotypic Plasticity, Predictive Modelling

## 1. Introduction

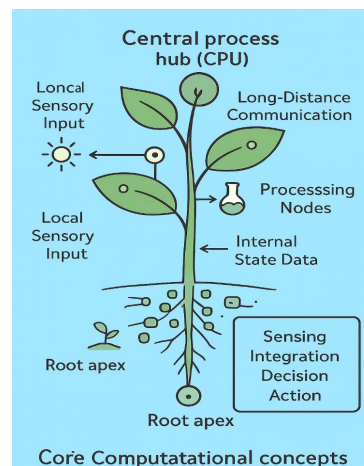
For centuries, plants were perceived as static, sessile life forms, their behaviors seen as simple, deterministic tropisms. This view is obsolete. A plant exists in a state of continuous, multi-input assessment, simultaneously monitoring light quality/quantity, temperature gradients, humidity, soil nutrient patches, mechanical forces, gravity, and the presence of pathogens or symbionts [1]. It must integrate this vast, noisy dataset with internal states—its developmental stage, metabolic status, and circadian rhythm—to execute appropriate, fitness-enhancing decisions: Where should the root grow? When should it flower? How should resources be allocated between defense and growth?

To achieve this, plants employ a sophisticated biological substrate for information processing. This article posits that the most fruitful framework for understanding these capabilities is to view the plant cell as a basic computational unit and the plant as a decentralized computing network [2]. In this paradigm:

- Sensors are membrane receptors (e.g., phytochromes, receptor kinases).
- Signal Transducers are second messengers ( $\text{Ca}^{2+}$ , ROS, inositol phosphates).
- Logic Gates & Processors are protein-interaction networks, phosphorylation cascades, and gene regulatory circuits.
- Memory is encoded in epigenetic marks, persistent protein modifications, and hormonal states.
- Wiring & Network Protocols are the vascular systems (xylem/phloem) and electrical signaling, transmitting processed information between distant processing nodes (e.g., root and shoot apices).

This computational perspective does not imply consciousness but highlights the existence of evolved, complex algorithms for survival. It bridges molecular biology with whole-plant physiology, offering a unified language to

describe phenomena like developmental plasticity, stress priming, and systemic acquired resistance [3]. This article will delineate the architecture of this cellular computation, explore the integration hubs within the plant body, and conclude with the transformative applications of this viewpoint for science and agriculture.



**Figure 1.** The plant as a distributed computational network.

Figure 1 presents a conceptual model of a plant as a distributed computational system, emphasizing how biological structures parallel elements of information processing. The shoot apical meristem and the root apex are highlighted as “Central Processing Hubs (CPUs),” representing regions where large amounts of sensory information are integrated and major developmental decisions are made. Surrounding tissues and individual cells function as “processing nodes,” collectively contributing to the plant’s ability to interpret and respond to its environment.

The diagram illustrates several categories of information flow. “Local Sensory Input” refers to environmental signals such as light, soil nutrients, water availability, and mechanical stimuli, which are detected directly by nearby cells. These inputs feed into internal pathways labeled “Internal State Data,” which capture the plant’s physiological status, often conveyed through hormones, metabolites, and other biochemical signals. “Long-Distance Communication” indicates the transport of information through vascular tissues, allowing distant organs—such as roots and shoots—to coordinate their reactions despite being physically separated.

At the bottom of the illustration, the foundational computational functions—sensing, integration, decision, and action—summarize how the plant processes information. First, the plant perceives environmental and internal cues; second, it integrates these signals across multiple tissues; third, it makes context-dependent decisions about growth, resource allocation, and stress responses; and finally, it executes these actions through physiological or morphological changes. Overall, the diagram conveys that plant behavior emerges not from a single “brain,” but from a distributed network of interacting nodes that collectively perform complex computation-like processes.

## 2. The Cellular Hardware: Sensors, Transducers, and Logic Circuits

Every plant cell is equipped with a toolkit for molecular computation.

### 2.1 Input Layer: Sensory Apparatus

Cells deploy an array of specialized receptors. Photoreceptors (phytochromes, cryptochromes) detect light direction, duration, and spectral composition. Mechanosensitive channels sense touch, gravity, and osmotic pressure. A vast family of receptor-like kinases (RLKs) perceives molecular patterns from pathogens (Microbe-Associated Molecular Patterns, MAMPs), cell wall integrity, and peptide hormones. This layer converts physical and chemical stimuli into primary biochemical signals [4].

### 2.2 Processing Core: Second Messengers and Kinase Networks

Primary signals are amplified and digitized into common intracellular currencies—second messengers like cytosolic calcium ( $\text{Ca}^{2+}$ ) waves, reactive oxygen species (ROS) bursts, and pH changes. These act as bit-like signals, where amplitude, frequency, and spatial localization encode information [5]. For example, a specific  $\text{Ca}^{2+}$  signature can distinguish between a touch stimulus and a pathogen attack.

These messengers activate downstream processors, primarily protein kinases and phosphatases. Cascades like the Mitogen-Activated Protein Kinase (MAPK) modules function as signal integrators and logic gates. A kinase can act as an AND gate, requiring phosphorylation from two upstream signals to become active, thereby only triggering a response when multiple conditions are met [6].

### 2.3 Memory and Output: Gene Regulatory Networks and Cellular Remodeling

The final output of cellular computation is often a change in gene expression. Transcription factors, themselves controlled by phosphorylation and other modifications, form complex gene regulatory networks (GRNs). These GRNs

can exhibit bistability, creating persistent "on" or "off" states that serve as cellular memory (e.g., vernalization). The output is a reprogrammed cell: altering its metabolism, wall properties, division plane, or fate [7].

### 3. Specialized Processing Hubs: The Root Apex and Shoot Apical Meristem

While all cells compute, certain tissues act as specialized command centers, integrating local and systemic data to direct organism-wide strategy.

#### 3.1 The Root Apex: A Sensory-Processor for Soil Exploration

The root tip is a premier example of an embedded environmental computer. Its structure-with the quiescent center, stem cell niche, and transitioning zones-is optimized for sensing and decision-making [8].

- **Sensory Module:** The root cap houses statocytes for gravity perception, cells sensitive to humidity, touch, and chemical gradients (nitrate, phosphate).
- **Integration & Decision:** The transition zone is proposed as a central processing site. Here, conflicting signals (e.g., high nitrate to the right, obstacle to the left) are integrated. The output is a directional instruction: adjust the rate and angle of cell elongation in the elongation zone via asymmetric auxin redistribution (a PIN-formatted auxin efflux) [9].
- **Algorithm in Action:** The classic "root foraging algorithm" involves oscillatory growth and preferential proliferation in nutrient-rich patches, a behavior reminiscent of an exploration-exploitation optimization strategy.

#### 3.2 The Shoot Apical Meristem (SAM): The Developmental Fate Scheduler

The SAM is the plant's architectural planning unit. It processes internal (hormonal, metabolic) and external (photoperiod, temperature) cues to make fate decisions for aerial organs [10].

- **Signal Integration:** The SAM integrates long-range signals like florigen (FT protein), which conveys photoperiod information from leaves, with local cues like cytokinin and auxin gradients.
- **Decision Logic:** Based on this integrated input, the meristem executes a developmental program. It decides whether to initiate a new leaf primordium, a branch, or a flower [11]. This involves complex spatial patterning, where cells interpret their position within a morphogen gradient (e.g., auxin maxima) to determine identity-a classic example of positional information processing.

### 4. Long-Distance Communication: The Plant's Network Protocol

Coordinated decision-making requires communication between distant hubs. Plants use both hydraulic and electrical signaling as their "network protocols."

#### 4.1 Rapid Electrical Signaling: Action Potentials and Variation Potentials

Triggered by wounding or severe stress, electrical impulses travel rapidly through the phloem and plasma membranes, akin to a network broadcast alert. These signals can quickly activate defense responses in distant leaves, preempting further attack [12].

#### 4.2 Systemic Chemical Signaling: Hormones as Information Packets

Phytohormones are the primary carriers of processed information. A stressed root does not send raw data; it synthesizes abscisic acid (ABA), a chemical summary of its "water deficit status." This ABA travels to shoots, where it is reinterpreted in the context of local conditions to execute stomatal closure. This represents a hierarchical information processing system.

### 5. System-Level Integration and Decision Making

The true computational power emerges from the interaction of all components. Plants perform system-level optimizations [13].

#### 5.1 Resource Allocation: A Continuous Cost-Benefit Analysis

Plants constantly balance carbon and nitrogen budgets between roots (acquisition), shoots (photosynthesis), and reproduction. This is governed by complex feedback loops involving sugars, nitrogen compounds, and hormones (e.g., cytokinin, auxin). The plant effectively runs a continuous linear programming model, maximizing growth under multiple constraints (light, water, nutrients).

#### 5.2 Predictive Behavior and Stress Priming

Plants exhibit anticipatory responses. Exposure to a mild stress can "prime" the defense system, leading to a faster, stronger response to a subsequent stress-a form of associative memory. Similarly, circadian rhythms allow plants to anticipate daily light/dark transitions, preparing their metabolism in advance [14]. This is computation extended over time.

### 5.3 Plant Computational Strategies in Ecological and Evolutionary Contexts

The computational capacities of plants have been shaped by natural selection over evolutionary time scales, reflecting their sessile lifestyle and the need to respond adaptively to environmental variability. From an ecological perspective, plant “decisions” are not made in isolation but are embedded within a dynamic network of interactions with competitors, mutualists, and herbivores [15]. For instance, when a plant is attacked by insects, it can emit volatile organic compounds that serve as warning signals to neighboring conspecifics and attract predators of the herbivores—a form of distributed communication protocol based on chemical signaling, illustrating collective-level computational coordination within plant populations.

From an evolutionary computation standpoint, plant populations can be viewed as parallel evolutionary systems, where each individual executes survival algorithms encoded in its genome and transmits successful strategies to offspring through reproduction. Phenotypic plasticity itself constitutes an adaptive algorithm at the evolutionary level, enabling plants to optimize performance under fluctuating conditions. For example, certain species adjust plant height and branching angles in high-density stands to maximize light capture—a behavior analogous to self-optimizing load balancing in distributed computing systems [16].

Moreover, plant-microbe symbiotic systems, such as mycorrhizal networks, form cross-species computational consortia. Mycelial networks not only transport nutrients but also relay defense signals and hormonal information between plants, effectively constructing an underground “internet of information.” This inter-organismal signal integration suggests that plants and their symbiotic microbiota can be regarded as an extended computational system, with information-processing capabilities that transcend the boundaries of a single organism.

This perspective also offers a novel lens through which to interpret major evolutionary innovations in plants. Key events—such as the emergence of vascular systems or the origin of floral organs—can be understood as major upgrades in biological computational architecture, enhancing the plant’s ability to process information and coordinate development in complex environments. Thus, the plant computation framework not only applies to physiological and developmental studies but also provides a unifying theoretical scaffold for macro-scale evolutionary ecology.

**Table 1.** Computational analogs in plant signal processing and decision-making.

Computational Concept	Plant Biological Component / Process	Example	Functional Role
Sensor / Input Device	Photoreceptors, Mechanoreceptors, Receptor Kinases	Phytochrome sensing red/far-red light; FLS2 detecting bacterial flagellin	Converts environmental or internal stimuli into a primary biochemical signal.
Signal Transducer / ADC	Second Messenger Systems (Ca <sup>2+</sup> , ROS, IP <sub>3</sub> )	Cytosolic Ca <sup>2+</sup> oscillations in response to touch or pathogen elicitors	Amplifies and converts analog sensory input into a digital-like, transmissible intracellular signal.
Logic Gate	Protein Kinase/Phosphatase Networks	MAPK cascade requiring sequential, conditional phosphorylation (AND gate logic).	Integrates multiple signals to produce a specific, conditional output. Enables decision nodes.
Processor / CPU	Gene Regulatory Networks (GRNs)	Network controlling floral organ identity (e.g., ABC model).	Executes complex developmental programs by integrating transcription factor activities. Determines cell fate.
Memory (RAM/HDD)	Epigenetic Marks (DNA methylation, histone mods); Persistent Protein States	Vernalization: cold exposure induces stable histone modifications that “remember” winter, promoting flowering.	Stores information about past experiences to influence future responses. Enables priming and adaptation.
System Bus / Network Protocol	Vascular System (Xylem/Phloem); Electrical Signaling	Transport of ABA (stress signal) via xylem; Rapid electrical wound signaling.	Transports information packets (hormones, RNAs) or rapid alerts between distant organs. Enables whole-plant coordination.
Central Processing Unit (CPU)	Root Apical Meristem; Shoot Apical Meristem (SAM)	Root apex integrating gravity, touch, and nitrate signals to direct growth; SAM integrating photoperiod (florigen) and hormonal cues to initiate flowering.	Specialized hubs for high-level integration of local and systemic data, issuing commands for growth or developmental transitions.
Algorithm	Phenotypic Plasticity Patterns; Foraging Behavior	Shade Avoidance Syndrome: algorithm integrating R:FR light ratio, neighbor proximity, and internal resources to decide stem elongation.	A pre-programmed, evolved set of rules for solving an ecological problem (e.g., optimizing light capture, nutrient foraging).
Cost-Benefit Model	Source-Sink Resource Allocation	Dynamic partitioning of photoassimilates between growing fruits, storage roots, and new leaves based on sink strength and vascular connectivity.	A continuous optimization process to maximize fitness under resource constraints.

Table 1 summarizes how plant signal processing and decision-making can be interpreted through the framework of computational concepts. Each row aligns a classical idea from computer science—such as sensors, logic gates, memory,

processors, algorithms, and cost-benefit models-with specific biological components or processes in plants. The table shows that many cellular and physiological mechanisms in plants perform functions analogous to those in engineered computational systems.

For example, “Sensors / Input Devices” correspond to plant photoreceptors, mechanoreceptors, and immune receptors, which detect environmental cues such as light, mechanical forces, or pathogen signals and convert them into biochemical information. “Signal Transducers / ADCs” include second-messenger systems like calcium and reactive oxygen species; these amplify raw stimuli and transform them into digital-like intracellular signals. “Logic Gates” are represented by kinase/phosphatase signaling networks, which integrate multiple inputs and produce conditional outputs much like AND or OR operations in electronics.

Higher-level computational roles are also mapped. Gene regulatory networks function as “Processors / CPUs,” because they integrate diverse information streams and control developmental programs. Epigenetic modifications and persistent molecular states serve as biological “Memory,” enabling plants to store information about past experiences and adjust future responses. The vascular system-through the movement of hormones, RNAs, and electrical signals-acts as the “System Bus,” coordinating communication between distant organs.

Finally, the table highlights how whole-plant behaviors resemble computational algorithms or optimization models. Phenotypic plasticity, such as shade avoidance or altered root growth, functions like an adaptive algorithm that adjusts growth to maximize fitness under changing conditions. Cost-benefit models such as source-sink resource allocation illustrate how plants continually optimize decisions about distributing nutrients and energy.

Overall, the chart demonstrates that plant biology can be interpreted through a computational lens, with biological structures and processes fulfilling roles analogous to engineered information-processing systems.

## **6. Implications and Future Perspectives: Engineering Computable Plants**

Viewing plants through a computational lens has profound implications.

### **6.1 For Basic Science**

It provides a unifying framework to design experiments and models. We can move from studying isolated pathways to reverse-engineering the algorithms of resilience. Questions shift from “What gene is induced?” to “What is the decision logic that triggers this response?”

### **6.2 For Predictive Agriculture**

This perspective enables *in silico* plant modeling with higher fidelity. By understanding the “code,” we can build digital twins of crops to simulate their responses to drought, disease, or novel management practices under climate scenarios.

### **6.3 For Crop Design**

The goal becomes engineering smarter crops. This doesn't just mean adding a single stress-tolerance gene, but potentially rewiring decision-logic circuits. Could we design roots with a more efficient “foraging algorithm”? Could we tune the flowering time decision network for greater climate robustness? Synthetic biology approaches could be used to install novel, logic-based sensor-response modules in plants.

### **6.4 Challenges, Controversies, and Future Directions: Refining the Plant Computation Framework**

While the computational system perspective offers a powerful heuristic framework, it is not without its conceptual challenges and controversies. Acknowledging these is essential for the paradigm's maturation.

**Conceptual Challenges and Avoiding Anthropomorphism:** A primary critique centers on the appropriateness of terms like “computation,” “decision,” and “intelligence.” Critics argue these risk unwarranted anthropomorphism, improperly attributing human cognitive properties to plants. A robust defense of this framework requires strict operational definitions: plant “computation” refers to unconscious, evolutionarily honed adaptive information processing; their “decisions” are the deterministic or probabilistic outputs of biochemical networks governed by evolved logic gates given specific inputs. The framework's strength lies in its value as a functional analogy and a formal modeling tool, not in postulating consciousness or intent.

**Quantification and Modeling Complexities:** Translating abstract computational analogies into testable, quantitative models is a formidable challenge. Plant signaling networks are characterized by high nonlinearity, redundancy, context-dependence, and noise. A single gene regulatory network may participate simultaneously in development, stress response, and metabolism, with its “logic function” dynamically shifting depending on cell type, developmental stage, and environmental history. Building truly predictive models of “plant algorithms” will require integrating multi-omics data with single-cell spatiotemporal dynamics and developing new computational theories capable of handling such complexity-a frontier at the intersection of systems biology and theoretical computer science.

**Beyond the Individual: Computation at Population and Ecosystem Scales:** The current framework primarily focuses on the individual plant. However, plants are nodes in larger networks, communicating via rhizosphere signaling, volatile emissions, and mycorrhizal networks to form extended information-processing systems. Future research must explore

how computation operates at this collective level: How is information encoded, transmitted, and integrated across different individuals? Are there emergent, population-level optimizations or "consensus" decisions? This would elevate plant computation theory from individual physiology to population and evolutionary ecology.

## 7. Conclusion

The evidence presented in this review compellingly supports a paradigm shift: the plant cell is a sophisticated computational unit, and the plant as a whole constitutes a distributed, adaptive biological computing network. Far from being passive responders, plants actively perceive a multifaceted environment, integrate noisy and often contradictory signals across spatial and temporal scales, access stored information from past experiences, and execute decisions that optimize fitness within ecological constraints. Framing plant biology through the lens of computation---borrowing rigorously from concepts in computer science, systems engineering, and information theory---transcends mere metaphor. It emerges as a powerful, predictive theoretical framework that unifies molecular mechanisms with whole-organism behavior and ecological strategy.

This computational perspective fundamentally reshapes our ontological understanding of plant intelligence. It moves the discourse beyond philosophical debates about consciousness and toward a functional understanding of evolved biological algorithms. Plants "compute" not with silicon, but with proteins, ions, and hormones; they "store memory" not in magnetic bits, but in epigenetic marks and persistent molecular states; they "communicate" not via Ethernet, but through vascular bundles and electrical currents. Recognizing these equivalences does not diminish the biological wonder but rather illuminates the convergent evolution of problem-solving strategies across living and engineered systems. It reveals that the principles of efficient information processing, robust decision-making under uncertainty, and decentralized network coordination are universal challenges, addressed through different substrates by evolution and human engineering alike.

The implications of this view are profound and tripartite. For basic science, it provides a generative framework for experimental design and modeling, urging a shift from studying isolated pathways toward reverse-engineering the complete algorithms of resilience and development. The key question evolves from "which gene is activated?" to "what is the *decision logic* that integrates these specific inputs to produce that adaptive output?" For applied agriculture, it enables the development of *in silico* digital twins of crops with unprecedented fidelity. By deciphering the plant's "operating code," we can simulate responses to complex, multi-stress scenarios predicted under climate change, revolutionizing predictive agronomy and sustainable management. For crop improvement, the ultimate frontier becomes engineering smarter plants. This extends beyond inserting single trait genes toward rationally rewiring endogenous decision-logic circuits---for instance, designing root systems with optimized foraging algorithms or reprogramming flowering time networks for greater environmental robustness using synthetic biology.

Looking forward, this paradigm opens interdisciplinary frontiers and reveals a deeper insight: life, at its most fundamental level, is a continuous process of information exchange. [Enhanced Conclusion]The algorithms honed in plants over eons represent robust, energy-efficient, and distributed solutions to Earth's environmental challenges. Future progress will advance along three axes. Theoretically, it demands deeper integration with complex systems science, information theory, and cognitive science to develop a formal "biological information processing" language that moves beyond analogy. Technologically, it inspires dual applications: plant-inspired computation for designing resilient algorithms and networks, and programmable plant synthetic biology, where genetic circuits are assembled like software to execute human-defined environmental logic. Ethically and conceptually, it fosters a renewed respect for plant capabilities, potentially reshaping agricultural ethics and our relationship with the biosphere.

Therefore, decoding the "living code" of plants is more than an analytical pursuit. It is an invitation to understand the biological algorithms of life itself. By doing so, we gain not only a deeper appreciation for the strategic intelligence of the vegetal world but also acquire the conceptual tools to predict, guide, and ultimately partner with plants in building a more resilient and sustainable future. The journey from viewing plants as static entities to recognizing them as dynamic, information-processing agents marks a significant leap in our scientific understanding---one that promises to bear fruit across fundamental biology, technological innovation, and global sustainability.

## References

- [1] Trewavas, A. (2017). The foundations of plant intelligence. *Interface Focus*, 7(3), 20160098. <https://doi.org/10.1098/rsfs.2016.0098>
- [2] Whippo, C. W., & Hangarter, R. P. (2006). Phototropism: bending towards enlightenment. *The Plant Cell*, 18(5), 1110-1119. <https://doi.org/10.1105/tpc.105.039669>
- [3] Monshausen, G. B., & Gilroy, S. (2009). Feeling green: mechanosensing in plants. *Trends in Cell Biology*, 19(5), 228-235. <https://doi.org/10.1016/j.tcb.2009.02.005>
- [4] Van Norman, J. M., et al. (2011). Intercellular communication during plant development. *The Plant Cell*, 23(3), 855-864. <https://doi.org/10.1105/tpc.111.082982>
- [5] Bassel, G. W. (2018). Information processing and distributed computation in plant organs. *Trends in Plant Science*, 23(11), 994-1005. <https://doi.org/10.1016/j.tplants.2018.08.006>
- [6] Forde, B. G., & Lorenzo, H. (2001). The nutritional control of root development. *Plant and Soil*, 232(1), 51-68. <https://doi.org/10.1023/A:1010329902165>

- [7] De Smet, I., et al. (2010). Bimodular auxin response controls organogenesis in *Arabidopsis*. *Proceedings of the National Academy of Sciences*, 107(6), 2705-2710. <https://doi.org/10.1073/pnas.0915001107>
- [8] Hedrich, R., & Neher, E. (2018). Venus flytrap: how an excitable, carnivorous plant works. *Trends in Plant Science*, 23(3), 220-234. <https://doi.org/10.1016/j.tplants.2017.12.004>
- [9] Choi, W. G., et al. (2016). Orchestrating rapid long-distance signaling in plants with Ca<sup>2+</sup>, ROS and electrical signals. *The Plant Journal*, 90(4), 698-707. <https://doi.org/10.1111/tpj.13492>
- [10] Kollist, H., et al. (2019). Rapid responses to abiotic stress: priming the landscape for the signal transduction network. *Trends in Plant Science*, 24(1), 25-37. <https://doi.org/10.1016/j.tplants.2018.10.003>
- [11] Mittler, R., et al. (2011). ROS signaling: the new wave? *Trends in Plant Science*, 16(6), 300-309. <https://doi.org/10.1016/j.tplants.2011.03.007>
- [12] Eveland, A. L., & Jackson, D. P. (2012). Sugars, signalling, and plant development. *Journal of Experimental Botany*, 63(9), 3367-3377. <https://doi.org/10.1093/jxb/err379>
- [13] Jaeger, K. E., & Wigge, P. A. (2007). FT protein acts as a long-range signal in *Arabidopsis*. *Current Biology*, 17(12), 1050-1054. <https://doi.org/10.1016/j.cub.2007.05.008>
- [14] Malamy, J. E., & Ryan, K. S. (2001). Environmental regulation of lateral root initiation in *Arabidopsis*. *Plant Physiology*, 127(3), 899-909. <https://doi.org/10.1104/pp.010406>
- [15] Silva-Navas, J., et al. (2016). D-Root: a system to cultivate plants with the root in darkness or under different light conditions. *The Plant Journal*, 84(1), 244-255. <https://doi.org/10.1111/tpj.12998>
- [16] Zhu, J. K. (2016). Abiotic stress signaling and responses in plants. *Cell*, 167(2), 313-324. <https://doi.org/10.1016/j.cell.2016.08.029>