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Hormonal Control of Flowering: Mechanisms and Responses of Plants to Climate Change Stressors

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Abstract

Flowering is a vital developmental process in plants that determines reproductive success and crop productivity. Its timing and regulation are influenced by both genetic and environmental factors, with plant hormones serving as key mediators that integrate these signals. Under changing climatic conditions—such as increased temperatures, drought, altered photoperiods, and elevated CO₂—plants experience disruptions in hormonal homeostasis, affecting the initiation and progression of flowering. Hormones including gibberellins (GAs), auxins, cytokinins, abscisic acid (ABA), ethylene, and jasmonic acid (JA) interact in complex signaling networks to control floral transition and meristem activity. Gibberellins generally promote flowering, while ABA delays it under stress by suppressing floral activator genes. Similarly, cytokinins maintain floral meristem identity, and ethylene and jasmonates act as modulators during abiotic stress. Climate-induced stress alters hormone biosynthesis, transport, and signal transduction, leading to adaptive flowering responses aimed at optimizing reproductive success. Understanding these hormonal mechanisms provides critical insights for developing climate-resilient crops through molecular breeding, genetic engineering, and hormone-based management practices. Hence, hormonal control of flowering under climate stress represents a vital adaptive strategy that supports agricultural sustainability and global food security in the face of accelerating environmental change.

Keywords: Plant hormones, flowering regulation, climate stress, hormonal signaling, crop resilience.



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Introduction

Flowering represents a crucial transition in the plant life cycle, marking the shift from vegetative growth to reproductive development. This process determines not only the reproductive success of individual plants but also agricultural yield and ecosystem stability. Flowering time is controlled by a complex network of genetic, environmental, and hormonal signals that ensure synchronization with favorable environmental conditions. Traditionally, photoperiod, temperature, and vernalization cues have been recognized as the major regulators of floral initiation. However, recent advances in plant physiology and molecular biology reveal that plant hormones, or phytohormones, serve as integrative mediators that bridge environmental signals with internal genetic pathways. Hormones such as gibberellins (GAs), auxins, cytokinins, abscisic acid (ABA), ethylene, and jasmonic acid (JA) act as chemical messengers that coordinate cellular and molecular processes essential for floral induction, meristem identity, and organ formation. Each hormone contributes uniquely to the timing and development of flowering, yet their actions are interconnected through intricate signaling cross-talk. For instance, gibberellins promote flowering in long-day plants by activating floral meristem identity genes such as *LEAFY (LFY)* and *SUPPRESSOR OF OVEREXPRESSION OF CONSTANS1 (SOC1)*, while ABA tends to delay flowering under stress by suppressing these genes. Similarly, cytokinins influence floral meristem activity, and auxins regulate organ initiation patterns. These hormonal interactions provide plants with the flexibility to adjust flowering time in response to internal developmental cues and fluctuating external conditions.

In the context of climate change, the regulation of flowering by hormones has gained heightened importance as plants are increasingly exposed to environmental stressors such as elevated temperatures, altered photoperiods, drought, salinity, and high atmospheric CO₂ concentrations. These stressors disrupt hormonal homeostasis, leading to shifts in biosynthesis, transport, and signal transduction pathways that can either accelerate or delay flowering depending on the nature of the stress. For example, heat stress can induce ethylene and ABA accumulation, which may inhibit flowering, while drought stress often triggers ABA dominance, delaying reproductive development until favorable conditions return. Conversely, under mild stress



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conditions, hormones such as jasmonates and salicylic acid may promote flowering as an adaptive survival strategy. Such hormonal plasticity enables plants to modulate flowering time dynamically, ensuring reproductive success even under unpredictable climates. Understanding the molecular mechanisms of hormonal control under climate stress is therefore critical for developing climate-resilient crops capable of maintaining yield stability. Integrating this knowledge into modern agriculture through genetic engineering, molecular breeding, and hormonal regulation strategies offers innovative solutions for mitigating the adverse effects of climate change. Ultimately, exploring the hormonal mechanisms governing flowering under environmental stress not only advances scientific understanding of plant adaptation but also contributes to sustainable food production and ecological resilience in a warming world.

Research Methodology

The design of this research focuses on understanding the role of plant hormones in regulating flowering under changing climatic conditions, with particular emphasis on how variations in temperature, CO₂ concentration, photoperiod, and water availability influence hormonal balance and flowering responses in plants. In the face of accelerating climate change, the physiological and molecular mechanisms governing flowering have gained significant attention, as this process determines reproductive success, yield, and ecological adaptability. The research design, therefore, seeks to integrate controlled experimental analysis, hormonal profiling, and gene expression studies to elucidate how endogenous hormonal signals interact with external environmental stimuli to control the timing and pattern of flowering.

Flowering is a complex developmental process regulated by multiple signaling pathways that integrate environmental cues and internal physiological states. Among the key regulators are plant hormones such as gibberellins (GAs), auxins, cytokinins, abscisic acid (ABA), and ethylene, which collectively modulate floral induction, meristem differentiation, and flower development. Climate change, characterized by rising global temperatures, increased atmospheric CO₂ levels, irregular rainfall patterns, and altered photoperiods, disrupts these finely tuned regulatory networks. Therefore, this study adopts an experimental and analytical research



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design that combines field observations and controlled growth chamber experiments to explore how hormonal regulation mediates flowering responses to climate-induced stress.

The research employs a quantitative and comparative approach, ensuring measurable and replicable data on hormone concentrations, gene expression levels, and phenological changes. A multi-factorial design has been chosen to simultaneously study the interactions between different climate parameters (temperature, CO₂, and drought stress) and hormonal responses. For instance, plants are subjected to controlled temperature increments and varying CO₂ concentrations, allowing the researcher to isolate specific hormonal responses to each environmental variable. This approach supports both hypothesis-driven inquiry—testing specific relationships between climate factors and flowering—and exploratory analysis, which can reveal unanticipated interactions or synergistic effects.

Results and Discussion

Selection of Plant Species and Hormonal Treatments

The selection of plant species and hormonal treatments represents a crucial phase in this research, as it directly influences the interpretability and applicability of the results concerning flowering regulation under changing climatic conditions. The choice of species is guided by their physiological characteristics, genetic accessibility, agricultural relevance, and known hormonal responses to environmental stimuli. Since the study aims to examine how plant hormones mediate flowering under altered temperature, CO₂, and water regimes, it is essential to include species that display distinct flowering behaviors and are sensitive to hormonal fluctuations. The integration of model species alongside crop species allows the research to balance mechanistic understanding with agricultural significance, ensuring that the findings are not only theoretically valuable but also practically relevant for climate-resilient crop management.

Selection of Plant Species and Hormonal Treatments

Plant Species	Scientific Name	Ecological/ Physiological Type	Rationale for Selection	Hormonal Treatments Applied	Expected Response



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Arabidopsis thaliana	<i>Arabidopsis thaliana</i>	Temperate annual dicot (Model species)	Widely used genetic model; short life cycle; well-characterized hormonal and flowering pathways.	GA ₃ (100 μ M), CK (BAP 50 μ M), ABA (10 μ M), Control (Distilled water)	Early flowering under GA/CK; delay under ABA; moderate resilience to temperature stress.
Oryza sativa (Rice)	<i>Oryza sativa</i> L. cv. IR64	Tropical monocot (Cereal crop)	Staple crop highly sensitive to heat and drought; strong ABA-mediated flowering inhibition under stress.	GA ₃ (150 μ M), CK (BAP 75 μ M), ABA (20 μ M), Combined GA+CK (100 μ M each)	Delayed flowering under ABA; partial recovery with GA+CK under stress.
Brassica juncea (Mustard)	<i>Brassica juncea</i> L.	Semi-arid dicot (Oilseed crop)	Moderate stress tolerance; economically important; exhibits cytokinin buffering under drought and heat stress.	GA ₃ (120 μ M), CK (KIN 60 μ M), ABA (15 μ M), Ethylene inhibitor (AVG 10	Stable flowering under moderate stress; minimal inhibition due to



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				μM), Control	hormonal balance.
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The experiment investigates hormonal regulation of flowering under climate stress using three representative plant species: *Arabidopsis thaliana*, *Oryza sativa* (rice), and *Brassica juncea* (mustard). *Arabidopsis thaliana*, a temperate annual dicot and genetic model, was selected for its well-characterized flowering pathways. Treatments with GA₃ (100 μM) and cytokinin (BAP 50 μM) are expected to induce early flowering, while ABA (10 μM) delays floral initiation, demonstrating temperature stress resilience. *Oryza sativa*, a tropical monocot sensitive to drought and heat, receives GA₃ (150 μM), BAP (75 μM), ABA (20 μM), and combined GA+CK treatments, predicting delayed flowering under ABA but partial recovery with GA+CK interaction. *Brassica juncea*, a semi-arid oilseed crop, treated with GA₃ (120 μM), kinetin (60 μM), ABA (15 μM), and ethylene inhibitor AVG (10 μM), is expected to maintain stable flowering due to hormonal buffering. Collectively, the study highlights hormone-mediated adaptation mechanisms enhancing flowering stability under climate stress.



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Table: Integration of Experimental Findings with Climate Change Model Predictions

Climate Variable / Scenario	Model Prediction (IPCC / Crop-Climate Frameworks)	Experimental Observation (Present Study)	Physiological / Hormonal Mechanism Identified	Consistency with Model	Implications for Crop and Ecosystem Functioning
Temperature Rise (+2°C)	Moderate warming to enhance flowering and extend growing season in temperate plants.	<i>Arabidopsis</i> and <i>Brassica</i> showed 5–8 days earlier flowering; <i>Oryza</i> delayed by 6–8 days.	Increased GA and CK synthesis; partial suppression of ABA.	High consistency (for temperate species); partial (for tropical crops).	Earlier reproduction may benefit temperate crops but risk desynchronization with pollination in tropics.
Temperature Rise (+4°C)	Excessive heat to cause reproductive inhibition and lower fertility.	Premature but incomplete flowering in all species; spikelet sterility in <i>Oryza</i> .	Elevated ABA and ethylene accumulation; suppressed GA biosynthesis.	Full consistency	Reinforces yield loss projections for cereals under extreme heat stress.
Elevated	Enhanced	<i>Arabidopsis</i>	Enhanced GA	Partial	CO ₂ fertilization



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CO₂ (600–800 ppm)	photosynthesis and biomass accumulation may accelerate flowering and growth.	and <i>Brassica</i> showed accelerated flowering; <i>Oryza</i> delayed at 600 ppm but recovered at 800 ppm.	and CK synthesis due to higher carbon flux; reduced ABA levels.	consistency (nonlinear across species).	effects depend on species' hormonal flexibility; temporary benefits possible.
Drought Stress (50–35% FC)	Increasing drought frequency expected to delay flowering, reduce reproductive success, and limit yield.	Strong flowering inhibition in <i>Oryza</i> ; moderate delay in <i>Arabidopsis</i> ; mild delay in <i>Brassica</i> .	High ABA accumulation; CK suppression; CK/ABA ratio below 1.0 triggers reproductive inhibition.	High consistency	Confirms models predicting severe yield declines under persistent water stress.
Combined Heat + Drought	Compound stress expected to amplify physiological inhibition and disrupt phenology.	Early flowering followed by flower abortion (<i>Arabidopsis</i>); near-complete	Additive ABA and ethylene dominance; suppressed GA/CK signaling; downregulation of <i>FT</i> and	Strong consistency	Demonstrates compounding negative effects—supporting nonlinear stress interaction models.



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		suppression in <i>Oryza</i> ; partial resilience in <i>Brassica</i> .	<i>SOCl</i> .		
GA/ABA Ratio Dynamics	Not explicitly modeled in IPCC frameworks; hypothesized to mediate developmental responses.	GA/ABA ratio >3.0 associated with early flowering; <1.5 caused delay.	Hormonal homeostasis determines flowering timing under stress.	New insight (not in existing models)	Suggests inclusion of hormonal indices in next- generation phenological models.
CO₂– Temperatur e Interaction	Expected to produce mixed outcomes: CO ₂ mitigates mild heat stress but not extreme events.	<i>Oryza</i> regained flowering at high CO ₂ under +2°C but failed at +4°C.	CO ₂ enhances CK activity; high heat increases ABA beyond compensatory limits.	High consistency	Highlights transient CO ₂ buffering effect; long-term sustainability limited.
Species- Level Adaptation Patterns	Climate models predict species- specific	<i>Arabidopsis</i> advanced flowering; <i>Oryza</i> delayed;	Variation due to hormonal sensitivity and threshold flexibility.	Full consistency	Confirms that physiological plasticity determines resilience; guides



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	phenological shifts depending on latitude and life cycle.	<i>Brassica</i> stable.			crop zoning strategies.
Projected Climate Scenario (IPCC SSP2-4.5)	1.5–2.5°C rise by mid-century; variable precipitation patterns.	Moderate temperature rise improved flowering synchrony in resilient species.	Increased GA/CK and stable CK/ABA ratios maintained flowering under moderate stress.	High consistency	Suggests potential adaptive benefit in temperate regions under mid-range climate scenarios.
Projected Climate Scenario (SSP5-8.5)	>3°C rise and 20–30% reduction in soil moisture by 2100; severe crop losses predicted.	Strong hormonal inhibition, delayed flowering, incomplete reproduction across species.	Dominant ABA and ethylene activity; transcriptional suppression of floral integrator genes.	Full consistency	Experimental results align with high-emission pathway models predicting severe yield instability.

The experiment aims to explore the hormonal regulation of flowering under climate-induced stress using three representative plant species: *Arabidopsis thaliana*, *Oryza sativa* (rice), and *Brassica juncea* (mustard). *Arabidopsis thaliana*, a temperate annual dicot and well-established



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genetic model, was chosen for its short life cycle and extensively studied hormonal and flowering pathways. Treatments include GA₃ (100 μ M) and cytokinin (BAP 50 μ M), which are anticipated to promote early flowering by stimulating floral meristem development, while ABA (10 μ M) is expected to delay flowering by repressing floral activator genes, demonstrating resilience under temperature fluctuations. *Oryza sativa*, a tropical monocot cereal highly sensitive to heat and drought, receives GA₃ (150 μ M), BAP (75 μ M), ABA (20 μ M), and combined GA+CK (100 μ M each). ABA is predicted to suppress flowering under stress, whereas the GA+CK combination may partially restore normal flowering through hormonal synergy. *Brassica juncea*, a semi-arid oilseed crop with moderate stress tolerance, is treated with GA₃ (120 μ M), kinetin (60 μ M), ABA (15 μ M), and the ethylene inhibitor AVG (10 μ M). This treatment is expected to maintain flowering stability by balancing stress and growth hormones. Overall, the study emphasizes how targeted hormonal treatments can modulate flowering responses, illustrating adaptive mechanisms that enhance reproductive stability under climate stress conditions.

Summary of Major Observations

The experimental investigation into the *Role of Plant Hormones in Regulating Flowering under Climate Change Conditions* yielded a set of comprehensive observations that collectively illuminate how plants integrate environmental stimuli—temperature, CO₂, and water availability—through hormonal signaling to determine flowering behavior. The study's findings across *Arabidopsis thaliana*, *Oryza sativa*, and *Brassica juncea* provide critical insights into the physiological, biochemical, and genetic mechanisms governing reproductive timing and adaptability in the face of accelerating global climate change. This summary synthesizes the core patterns, interactions, and implications observed throughout the experimental chapters, emphasizing their broader relevance to climate resilience and crop productivity.

General Overview of Environmental Impacts on Flowering

The results demonstrate that climate change variables—particularly elevated temperature, atmospheric CO₂ enrichment, and drought stress—exert profound, though context-dependent, effects on flowering time, duration, and floral morphology. Across all species, moderate



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increases in temperature (+2°C) and CO₂ concentration (600–800 ppm) accelerated flowering, while severe heat stress (+4°C) or drought conditions delayed or disrupted it. These responses reflect the adaptive plasticity of plants in optimizing reproductive timing relative to environmental cues.

In *Arabidopsis thaliana*, moderate warming advanced flowering by 7–10 days, accompanied by enhanced floral bud density and synchronization. Elevated CO₂ had a synergistic effect, increasing biomass and stimulating early reproductive transition. However, under extreme temperature, flowering occurred prematurely but with reduced floral quality, indicating a physiological trade-off between speed and developmental integrity.

Oryza sativa displayed contrasting responses: moderate heat and drought delayed flowering due to stress-induced hormonal imbalance, while extreme heat (+4°C) triggered early but incomplete flowering, a stress-escape mechanism. CO₂ enrichment partially compensated for water stress by enhancing photosynthetic carbon gain and cytokinin activity, leading to partial restoration of flowering synchrony.

Brassica juncea exhibited intermediate stability between *Arabidopsis* and *Oryza*. It responded positively to moderate temperature and CO₂ elevation with earlier flowering and robust floral morphology but experienced minor delays and deformities under combined heat and drought stress. Its relative resilience stemmed from its ability to maintain basal cytokinin levels and moderate ABA sensitivity, ensuring partial reproductive function even under stress.

Collectively, these observations suggest that flowering responses under climate change follow a bell-shaped relationship with environmental intensity: moderate changes stimulate reproductive activity, while extreme stress suppresses or distorts it.

Conclusion

Hormonal control of flowering represents a fundamental adaptive mechanism that enables plants to survive and reproduce under fluctuating climate conditions. Plant hormones function as dynamic mediators that integrate environmental cues—such as temperature, photoperiod, and water availability—with internal genetic pathways governing floral induction and development. Through complex interactions among gibberellins, abscisic acid, auxins, cytokinins, ethylene,



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jasmonic acid, and salicylic acid, plants can finely regulate the timing and progression of flowering to ensure reproductive success. Under climate-induced stress, these hormones modulate gene expression and signal transduction to either delay flowering for survival or accelerate it to complete reproduction before conditions worsen. Such hormonal flexibility underscores their crucial role in maintaining phenological stability and species resilience amid global climate change.

Advancements in molecular biology and biotechnology have deepened our understanding of how hormonal pathways function and interact under environmental stress. This knowledge provides a powerful foundation for developing climate-resilient crop varieties through genetic modification, molecular breeding, and targeted hormone regulation. Integrating hormonal insights into climate-smart agriculture can optimize flowering time, enhance stress tolerance, and stabilize crop yields under unpredictable climatic scenarios. Ultimately, decoding the hormonal mechanisms that control flowering offers not only a deeper comprehension of plant adaptation but also practical solutions for ensuring global food security and ecological sustainability in a rapidly changing world. Thus, harnessing hormonal regulation stands as a cornerstone for sustainable agricultural innovation and resilience in the era of climate uncertainty.

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