

Integrating Leaf and Canopy Photosynthesis Modelling for Enhanced Crop Improvement

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Abstract—Predicting crop yield by integrating molecular basis for crop improvement attracts increasing interest but remains a great challenge due to complex biological and physiological processes across various scales. Crop physiological models have been developed and successfully used in predicting crop growth, development, and grain yield formation in response to various environments for more than half century. The crop model realized a chain of non-linear processes from canopy development, canopy photosynthesis, biomass growth and allocation, grain yield formation, and nitrogen, water, and energy balance based on detailed physiological and agronomic experiments. On the other hand, leaf biochemical photosynthesis was also modelled with relatively simple equations four decades ago, which receives wide attention due to underlying explicit biochemical mechanisms. Apart from that, the model is based on photosynthetic events occurring on leaf cellular chloroplast, which more closely represents the fundamental molecular regulation. Molecular regulatory network of photosynthesis is made clear, but such achievement in improving grain yield has not been translated well in that the molecular basis is not directly reflected in whole plant performance and lacking predicting ability from cellular level to whole plant level. Consequently, connecting both types of models will greatly accelerate the efforts of realizing the prediction of molecular outcomes in grain yield formation. Here we reviewed development of leaf and canopy photosynthesis models, and discussed the connection of both types of models to support crop improvement.

Index Terms—Crop improvement, canopy photosynthesis, leaf biochemical photosynthesis, crop models, model connection, molecular basis, whole plant physiology

I. INTRODUCTION

A. Mission Switch in Crop Improvement

CROP yield depends on resource capture by plant, resource use efficiency, and harvest index. While extensive efforts to optimize resource capture and harvest index have brought crop productivity to a plateau with narrow margins for further enhancement [1], our grasp of the physiological and molecular mechanisms regulating resource

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dynamics—including capture, flow, and utilization—has advanced profoundly. Therefore, shifting the focus toward elevating resource use efficiency represents a promising new frontier for driving advancements in agricultural productivity [2].

B. Photosynthesis Offers Potentials for Crop Improvement

Through the process of photosynthesis, leaves intercept solar radiation and convert it into chemical energy by transforming carbon dioxide into stored carbohydrates. As a fundamental cellular biochemical pathway, leaf photosynthesis serves as the primary stage where critical resources (i.e., light, water, and nitrogen) are integrated, making it a pivotal target for optimizing resource use efficiency. Carbon assimilation within the leaf acts as the foundational energy source that drives crop growth, development, and yield formation. Research has demonstrated that leaf photosynthesis can be significantly boosted either by elevating CO₂ concentration or by optimizing the performance of key photosynthetic enzymes [3, 4]. Consequently, optimizing the biochemical activities of the photosynthetic process presents a significant opportunity for elevating crop productivity [5–8].

Enhancing photosynthetic capacity is considered a pivotal strategy for boosting agricultural productivity to meet the demands of expanding populations [5]. The clarification of photosynthetic biochemical pathways [9] boosts the currently processes of photosynthesis improvement. Since CO₂ is the raw material for carbon fixation, minimizing diffusion barriers is a top priority, specifically through the regulation of stomatal [10] and mesophyll conductance [11]. Under field conditions, the photosynthetic capacity of leaves is typically capped by electron transport limitations or the ability to harness the produced chemical energy [5]. In the context of light reactions, potential enhancements include extending the usable spectrum into the near infrared radiation (NIR) [12], optimizing canopy light distribution by reducing antenna sizes in top leaves [13, 14], and facilitating a faster relaxation of photoprotective heat dissipation at photosystem II (PSII) [15–17]. When light is non-limiting, carbon assimilation is often co-limited by the efficacy of the carboxylation step and the replenishment of its substrate [5].

Downstream processes in the dark reactions offer multiple targets for optimization, such as engineering C4 traits into C3 species [8], incorporating inorganic carbon-concentrating mechanisms from algae or cyanobacteria [18, 19], redesigning rubisco for current CO₂ level [20], and synthesizing photorespiratory bypasses [21, 22]. Furthermore, enhancing ribulose biphosphate (RuBP) regeneration [23] and improv-

ing light penetration to the lower canopy [24, 25] can also boost performance. While certain strategies currently appear more feasible, there is insufficient evidence to prioritize one specific method, as many of these targets have yet to test through field-based validation.

C. Limited Predictive Ability from Molecular and Biochemical Basis to Crop Yield

Crop yield does not always mirror the molecular drivers of leaf photosynthesis, as numerous non-linear processes govern the transition between cellular biochemical activities and yield. Consequently, efficient modeling tools are indispensable for evaluating and predicting the impact of molecular-scale modifications on whole-plant performance [26–28]. Additionally, the scientific consensus among physiologists is that crop yield represents a highly complex phenotype, where leaf-level carbon fixation is only one of many contributing physiological factors [8, 27]. This suggests that future breeding efforts should focus on identifying diverse trait variations and integrating them in a synergistic manner to achieve substantial yield gains.

D. Existing Basis for Modelling Prediction

Since 1965, canopy photosynthesis modelling has evolved into a comprehensive system capable of integrating environmental and management-related factors [29]. As the fundamental engine for plant growth and development within crop models, the underlying algorithms for canopy photosynthesis have shifted from empirical light-response curves [29] or biomass-light efficiency correlations [30] toward more mechanistic biochemical frameworks at the leaf level [28, 31]. These biochemical models offer a high-fidelity simulation of plant-environment dynamics, bridging the gap between molecular and multi-organismic scales [5].

With current scientific insights, it is now feasible to construct virtual crop canopies [26, 32]. Such models function as testing grounds for hypotheses aimed at maximizing carbon sequestration and applying optimization strategies for improved efficiency. To date, numerous mechanistic modules have been refined, encompassing gene regulation [32], photosynthetic biochemistry [33], mesophyll-cell diffusion dynamics [34], stomatal behavior [10], respiration [22, 35], and complex 3D microclimates within the canopy [36]. Despite their utility in identifying specific pathways for improvement, these isolated frameworks often fail to account for the interactions occurring within the entire plant system or the surrounding environment. The integration of these diverse modules into a multi-scale modeling framework can facilitate more accurate, system-wide predictions of photosynthetic performance across different biological levels.

E. Objectives of This Literature Review

To this end, this study first reviews the evolution of both canopy-level and biochemical-scale photosynthesis simulations. Subsequently, it evaluates the adaptability of these models in light of emerging scientific missions and constraints. Finally, we elucidate potential methods for synthesizing these two modeling paradigms to provide a robust theoretical foundation for crop improvement initiatives.

II. MODELLING CROP CANOPY PHOTOSYNTHESIS

Defined as the aggregate photosynthetic activity of all leaves in a stand, canopy photosynthesis captures the dynamic interactions occurring among plants and their external surroundings at the population scale, which functions as a foundational metric for estimating agricultural productivity [28] and plays a significant role in modulating global climate patterns by regulating atmospheric CO₂ levels [37]. The light dependence of a canopy deviates significantly from that of a single leaf; thus, canopy photosynthesis cannot be accurately estimated by merely aggregating the photosynthetic rates of all leaves [38]. Within the canopy, the carbon assimilation capacity of individual leaves is dictated by their placement, physical form, and physiological status [39]. Environmental variables inside the canopy, including the intensity and spectral composition of light, temperature, moisture levels, and CO₂ gradients, display pronounced spatiotemporal variability [36, 40, 41]. In addition to structural differences, leaves exhibit marked physiological divergence across the canopy profile [7]. Specifically, upper canopy leaves are characterized by increased rubisco content and leaf thickness, as well as higher chlorophyll a to chlorophyll b ratios, whereas lower leaves show different protein and pigment compositions [13, 42]. It is essential to recognize that leaves undergo developmental acclimation to their local light environment over time. Such adaptive plasticity in photosynthetic metabolism is particularly evident in shade-tolerant plants responding to intermittent light patches [43]. Despite the abundance of existing canopy models, only a limited number of them incorporate the intricate physiological variations and specific performance of each leaf. Accounting for these factors is necessary to ensure that modeled canopy gas exchange remains consistent with measured data.

A. Maximum Productivity Model

As the most rudimentary approach to simulating canopy net primary productivity (NPP), the maximum productivity model calculates output by adjusting a crop-specific potential maximum. This adjustment accounts for limiting environmental stressors such as insufficient light, sub-optimal temperatures, and the scarcity of water or nutrients [44].

The CENTURY model, frequently employed for simulating soil organic matter dynamics, adopts this simplified logic [44]. Given it focuses on underground carbon processes, the model avoids complex photosynthetic architectures [45]. However, this trade-off prevents the simulation of crop responses to varying levels of incident light. When applied at the site level, the model's reliability has been verified across diverse ecosystems, ranging from grasslands [46] to forests [47]. Findings suggest a satisfactory performance, with deviations typically remaining within $\pm 25\%$ of empirical measurements [48]. Nevertheless, because the datasets for calibration and validation frequently overlap [46], these results may not constitute a truly stringent assessment of the model's predictive accuracy. Similar to the maximum productivity approach, several other frameworks utilize coarse-grained techniques for canopy-scale simulations. Notable examples include terrestrial ecosystem model (TEM) [49], PnET [50], and integrated biosphere simu-

lator (IBIS) [51], all of which are extensively applied to study productivity within terrestrial ecosystems.

B. Radiation Use Efficiency (RUE) Model

Rooted in early efforts to correlate crop weight with solar radiation [29], the RUE model emerged from the documented linear relationship between cumulative intercepted light and dry matter production [52]. This concept has since been consolidated through extensive experimental and theoretical verifications [53]. Consequently, RUE is the biomass generated per unit of total radiation or photosynthetically active radiation (PAR), and has become a pivotal metric in productivity analysis and a core of crop growth simulation. In practice, canopy productivity is estimated by taking the product of the light intercepted (derived from light distribution models [54]) and a predefined RUE [55]. The RUE can be obtained by fitting a linear relationship between cumulative biomass accumulation and cumulative radiation interception.

Thereafter RUE has quickly received much interest and many studies carried out in determining RUE values in various crop species and environments. Variations in RUE are primarily determined by crop type, with C4 plants exhibiting the highest capacity, followed by non-legumes and legumes [55]. Furthermore, RUE is not a static constant; it is sensitive to nitrogen status [56] and light quality. Evidence suggests that a shift toward diffuse radiation can boost RUE by up to 50% in various species [57, 58]. The widespread adoption of RUE in diverse crop models is primarily attributed to its straightforward formulation and the relative ease of obtaining its governing parameters. A theoretical challenge within the RUE approach involves the mechanism by which non-linear leaf-level light responses are effectively scaled into a linear canopy-level representation. Evidence indicates that this linearity is an emergent property applicable only at two weeks or longer intervals, whereas the coupling between intercepted radiation and productivity remains non-linear on a short-term scale [59, 60]. Furthermore, synergistic interactions among multiple resource constraints contribute to the observed long-term linearity of light-limited canopy photosynthesis [61]. For instance, under nitrogen-deficient conditions, RUE is positively with leaf nitrogen concentration [56] and is sensitive to the vertical distribution of nitrogen within the canopy profile [62]. RUE values for maize, soybean, and grass have been reported to rise by 33%–50% when the light environment shifts toward lower total incident radiation but higher diffuse radiation [57, 58]. Echoing the sensitivities of leaf photosynthesis, RUE fluctuates in response to temperature, CO₂ concentration, and water availability [28]. Therefore, RUE-based models typically employ corrective functions to account for these environmental constraints and improve predictive robustness [63].

The aforementioned methods represent simplified frameworks for estimating canopy photosynthesis. Both the maximum productivity and RUE models were conceptualized through a “top-down” lens, grounded in empirical observations at the macroscopic scale, with their linkages to microscopic processes being explored only retrospectively. Consequently, these models necessitate a vast array of empirical

coefficients to encapsulate the influences of fluctuating environments and intrinsic plant physiology. This heavy reliance on empirical fitting serves as a major constraint on their predictive precision. Although effective for assessing photosynthetic potential within specific conditions, these approaches lack the robustness required for extrapolation to novel environments. Their primary deficiency lies in an inability to resolve the intricate interactions between canopy photosynthesis and external drivers, such as temperature, humidity, or CO₂ concentrations [26, 64]. As a result, such modeling paradigms may prove inadequate when the objective is to evaluate the consequences of physiological modifications or rapidly changing climatic scenarios.

C. Multi-Layer Approach

The inception and advancement of empirical frameworks for canopy photosynthesis were primarily driven by the formulation of mathematical leaf light-response curves. Concurrently, the integration of light attenuation models within the canopy significantly catalyzed the growth of this field [54]. The multi-layer framework operates by partitioning the whole canopy into numerous horizontal layers. Early versions of these models utilized exponential functions, such as the Beer-Lambert law, to simulate vertical light gradients. The extinction coefficient (k) is determined for different canopies primarily based on leaf inclination angles [54]. A key simplifying assumption of this approach is that all leaves within a given horizontal layer experience uniform light availability. The light-response behavior of individual leaves was typically represented by a rectangular hyperbola function. The canopy photosynthesis is the sum of individual leaf photosynthesis rates. This framework proved effective in providing a formal mathematical description of the essential drivers of canopy photosynthesis.

A subsequent development was a simplified multi-layer canopy model [65] that focused on vertical variations in wind speed and irradiance. However, it treated other environmental factors as uniform across whole canopy, including temperature, humidity, and CO₂ concentration. Almost at the same time, an advanced multi-layer framework was introduced [66] with expanded capabilities. It facilitates the calculation of real-time CO₂, heat, and radiation, allowing canopy photosynthesis to be estimated through empirical models of light absorption. Despite their inherent complexity, full multi-layer models serve as powerful tools for both modeling energy and mass exchange between canopy and the atmosphere and elucidating the underlying physiological and physical mechanisms.

Early modeling efforts assumed leaf homogeneity across the canopy due to limited computing resources. To incorporate vertical physiological gradients, researchers began by calculating leaf nitrogen profiles within the canopy and linking them to key light-response parameters [67]. While initial leaf photosynthesis model relied solely on light as a driving variable, the emergence of biochemical modeling [33] allowed for the simultaneous consideration of CO₂ and temperature. The subsequent addition of stomatal conductance functions [68] created a synergistic approach for predicting leaf-level gas exchange in varying environmental conditions [69, 70].

D. Big-Leaf Approach

The big-leaf model conceptualizes the entire canopy as a unified, single-layer leaf. This approach is predicated on the hypothesis that the spatial allocation of photosynthetic capacity throughout the canopy mirrors the gradient of light intensity [71]. Such a premise facilitates the direct scaling up of leaf-level light-response functions to the canopy scale. Additionally, it operates under the simplification that environmental variables, such as temperature and humidity, remain uniform across whole canopy. Consequently, the physiological reactions of the canopy to external drivers can be simulated via mechanistic biochemical models of leaf photosynthesis. This logic was pioneered by Farquhar [72], who stated that if leaf photosynthetic capacity is distributed proportionally to absorbed radiation and leaf-level response curves are consistent, the mathematical expressions for canopy carbon fixation retain the same structure as those for single leaves. This theoretical framework was promptly adopted and expanded within the domain of canopy photosynthesis research [71, 73].

Following this theory, various researchers have explored whether a vertical, exponential decline in leaf photosynthetic capacity occurs in tandem with radiation attenuation, following the principles of Beer's law. Given the robust linear correlation between leaf photosynthesis and leaf nitrogen levels [74], empirical validation has largely centered on the spatial partitioning of nitrogen within the canopy profile. Findings from optimization analyses demonstrate that nitrogen distribution in proportion to the light extinction represents a theoretical optimum for maximizing canopy carbon gain [71, 75]. It has been widely documented that leaf nitrogen and its corresponding photosynthetic capacity exhibit a marked downward trend along the vertical canopy axis [67, 76]. Model-based comparisons revealed that when total canopy nitrogen is conserved, the optimized nitrogen distribution closely mirrors observed patterns, suggesting that plants naturally tend toward this high-efficiency allocation strategy [67, 74].

Although the premise of a proportional reduction in photosynthetic capacity alongside light attenuation simplifies the integration of leaf-level models into canopy level, historical application has shown that it has not simplified integration. For example, the HYBRID model fails to account for diurnal fluctuations in canopy photosynthesis, relying instead on daily mean values for both irradiance and carbon fixation [77]. Consequently, this method disregards the critical interplay between the non-linear dynamics of leaf-level response models and the temporal variability of light absorption within the canopy structure. Nevertheless, even without internal parameter recalibration, the HYBRID model's efficacy was evaluated against global primary productivity and carbon stocks [78]. While the model exhibited satisfactory overall performance, certain inconsistencies with observations underscored specific domains necessitating refinement.

The big-leaf framework was later refined to include diurnal variations in solar radiation [79], leading to its successful implementation in models such as PROMOD [80] and CenW [81]. These models have undergone rigorous field-scale vali-

ation. For instance, PROMOD's predictive accuracy regarding annual increments was confirmed across 28 sites [80], while CenW was validated using experimental data on leaf nitrogen, soil moisture, and biomass partitioning from irrigation and fertilization trials [81]. A primary merit of the big-leaf approach is its compatibility with remote sensing data for light interception, which bypasses the complexity of simulating radiation transfer. This feature is leveraged by the simple biosphere model, version 2 (SiB2), which serves as the terrestrial surface module for the global climate model (GCM) [82]. Consequently, the model must function at high temporal resolutions (minutes) on a global scale. At such scales, incorporating the non-linear relationship between intercepted radiation and canopy photosynthesis becomes essential, making the big-leaf approach an appropriate modeling selection.

As previously noted, this approach is predicated on the assumption of a parallel exponential decline in both light intensity and leaf photosynthetic capacity within the canopy. Although this premise is technically an oversimplification, its practical application typically avoids introducing severe errors. A critical flaw is that leaf within any given layer comprises both sun and shaded leaves, each subjected to distinct light intensities [83]. Given the high temporal variability of sunlit leaves, the capacity for photosynthesis cannot be considered a simple linear function of the instantaneous light intensity. It is therefore imperative to partition the canopy into sunlit and shaded fractions, calculating their photosynthetic contributions independently. Empirical evidence from eddy covariance studies further indicates that canopies react differently to diffuse versus direct solar fluxes, necessitating a different modeling approach [84]. Both the heterogeneity of leaf exposure and the nature of incident light are explicitly accounted for in sun-shade modeling framework.

E. Sun-Shade Leaf Approach

The sun-shade leaf framework provides a comprehensive accounting of the distinct behaviors of direct and diffuse radiation, as well as the pervasive nature of diffuse flux within the canopy. Specifically, this modeling paradigm partitions the canopy into several layers, each of which is bifurcated into two distinct physiological populations: sunlit and shaded leaves. Within this approach, sunlit leaves are modeled as intercepting a combination of direct and diffuse radiation, whereas shaded leaves are restricted to diffuse-only interception. A foundational assumption of this approach is the spatial homogeneity of both radiation interception and photosynthetic activity within each leaf fraction [83]. By synthesizing the characterization of shaded leaves with the theory of optimal canopy nitrogen partitioning, a highly precise canopy photosynthesis model is realized [83, 85]. Building upon these premises, the sun-shade model [83] integrates a mechanistic biochemical model of leaf photosynthesis [33] to compute canopy-level CO₂ assimilation rates [33].

While the sun-shade model exhibits greater architectural complexity than the big-leaf approach, it maintains a mechanistic framework that does not necessitate additional parametric input. Its primary advantage is the strategic weighting of sunlit versus shaded leaf contributions, achieved by dividing

the canopy into these two distinct physiological classes. Consequently, this model offers superior predictive accuracy relative to the previously discussed empirical and simplified frameworks. A notable constraint, however, is its incompatibility with radiation interception data derived from remote sensing, thereby demanding more comprehensive information for robust parameterization.

The initial implementation of the sun-shade leaf approach was featured in the generic decomposition and yield (G'DAY) model [86], utilizing photosynthetic parameters derived directly from in field measurements. Despite its foundational role, the initial scalability of this model remained limited, as it lacked extensive validation and application across broader geographical contexts. The adoption of the sun-shade leaf approach within G'DAY was primarily motivated by the facility with which leaf-level responses to nitrogen, water availability, and temperature could be integrated into the canopy framework. This represented a significant advancement over earlier iterations of the G'DAY model, which relied on the RUE paradigm [87]. A prominent example of upscaling the sun-shade leaf approach for regional net primary productivity simulations is the CSIRO biospheric model (CBM) [85]. Functioning at a 30 min temporal resolution, CBM provides high-frequency simulations of gas and energy exchanges within the soil-plant-atmosphere continuum. Site-specific validations using wheat eddy covariance data have demonstrated its predictive accuracy [85], leveraging established physiological parameters to refine the light-response curvature [88].

F. 3D Modeling Approach

Integrated as the driving force within crop simulation systems, canopy photosynthesis models quantify carbon fixation throughout the growing season [29]. These range from simple linear maximum productivity models and RUE models [52, 53] (e.g., light interception and utilization simulator (LINTUL) [59], agricultural production systems simulator (APSIM) [89], and decision support system for agrotechnology transfer (DSSAT) [90]) to more complex leaf-aggregation approaches like the multi-layer models, big-leaf models, and sun-shade leaf models [29, 91, 92] (e.g., simple and universal crop growth simulator (SUCROS) [93] and genotype-by-environment interaction on crop growth simulator (GECROS) [94]). While certain models like APSIM allow for algorithmic flexibility [95], most share a common limitation: an overly simplify canopy geometry. This lack of architectural detail precludes an accurate representation of internal environmental heterogeneity, rendering such models inadequate for evaluating structural modifications or identifying breeding targets for enhanced light interception.

Recent advancements in functional-structural plant models (FSPMs) have enabled the characterization of intricate spatiotemporal dynamics regarding plant growth within a 3D context. These provide a robust foundation for the high-fidelity simulation of light distribution within complex canopy architectures. Diverging from traditional approaches that rely on simplified parameters such as extinction coefficient or leaf area index (LAI), this 3D framework reconstructs the canopy

by integrating granular architectural traits, including tiller and leaf counts, spatial positioning, and leaf orientation, with specific agronomic variables like planting density and row direction [96]. Subsequently, a ray-tracing algorithm is deployed, incorporating data on incident irradiance intensity, spectral composition, and angular directionality to compute the precise light intensity received by individual leaves within the canopy. By synthesizing these radiation inputs with leaf photosynthesis and stomatal conductance parameters, a comprehensive 3D canopy photosynthesis model is established. This framework serves as a versatile biophysical platform for the precise estimation of canopy photosynthesis. To empirically test the model's accuracy, a specialized canopy gas exchange measurement system was designed to provide direct observations of photosynthesis rates [97]. Comparative analyses demonstrated strong congruence between the simulated outputs and the observed data, confirming the model's predictive reliability.

A comparative study on canopy photosynthesis models with different dimensions demonstrated that crop yield predictions decreased by 12.0%–48.5% with increasing the resolution of light simulation [98]. However, the efficacy of the 3D canopy photosynthesis model depends on precise parameterization. Key requirements for this process include: (1) leaf-level biochemical and physiological parameters, such as maximum rate of rubisco carboxylation capacity and RuBP regeneration; (2) detailed architectural traits, such as tiller and leaf counts, leaf phyllotaxy, and leaf angle and orientation; and (3) agronomic configurations, including row spacing, plant density, etc. [96, 99]. These datasets can be acquired via traditional manual measurements or through advanced high-throughput phenotyping platforms. While manual characterization remains a labor-intensive and time-consuming bottleneck, it currently provides a baseline for model calibration. The rapid evolution of plant phenotyping technology is expected to mitigate the challenges of model parameterization. The 3D canopy photosynthesis model can provide more detailed structural and functional attributes of plants compared to other models; however, it entails greater effort in parameterization and is computationally expensive to execute [64]. Consequently, the selection between modeling approaches should be guided by the research objective: while RUE-based models are suitable for regional yield estimations due to their simplicity, 3D models are essential for identifying fine-scale breeding targets. As the technical barriers diminish in advanced genomics, phenomics, and computational, the 3D modeling approach will become an increasingly indispensable tool for crop improvement and understanding complex plant-environment interactions [100, 101].

The conceptual derivation of these canopy photosynthesis models follows two distinct paradigms. The big-leaf, multi-layer, sun-shade, and 3D modeling frameworks are rooted in a “bottom-up” approach, integrating fine-scale physiological non-linearities and their interactions with environmental drivers. Conversely, the maximum productivity and RUE methodologies adopt a “top-down” logic, derived initially from empirical measurements at the canopy scale.

Current canopy photosynthesis models demonstrate a high

degree of biophysical realism, accurately capturing land-atmosphere exchange rates for carbon, water, and energy. Although these models range from mechanistic to simplified, they maintain remarkable predictive precision. The process of scaling up from leaf to canopy is fundamental to elucidating the mechanisms governing fluctuations in gas and energy exchange. Furthermore, these models are indispensable for projecting the long-term responses of terrestrial ecosystem functions to climate change. However, robust predictions depend on high-resolution data regarding plant architectural and functional traits. As the plasticity of these traits in response to changing environments remains partially obscured, current projections are often characterized by substantial uncertainties. Nevertheless, the synergy between expanding flux networks and improved data-model assimilation techniques is paving the way for more robust and sophisticated global carbon cycle prediction.

III. MODELLING BIOCHEMICAL PROCESSES OF LEAF PHOTOSYNTHESIS

Theoretical insights into leaf photosynthesis have culminated in the development of the Farquhar-von Caemmerer-Berry (FvCB) model [33], a cornerstone of C3 plant physiology. By mathematically representing the biochemical pathways within the chloroplast, the model estimates net CO₂ uptake A by identifying the primary limiting factor at any given moment. Specifically, it calculates A as the lower value between the rubisco-limited assimilation rate A_c and the electron transport-limited rate A_j .

$$A = \min\{A_c, A_j\} \quad (1)$$

A_c is calculated as a function of the maximum carboxylation capacity of rubisco V_{cmax} by Eq. (2).

$$A_c = \frac{(C_c - \Gamma_*)V_{\text{cmax}}}{C_c + K_c(1 + O_c/K_o)} - R_d \quad (2)$$

where C_c is the CO₂ partial pressure at the carboxylating sites of rubisco. K_c and K_o are Michaelis-Menten constants of rubisco for CO₂ and O₂, respectively. Γ_* is the CO₂ compensation point in the absence of day respiration R_d .

In the calculation of A_j , the FvCB model assumes 100% non-cyclic e-transport, thus excluding cyclic e-transport around photosystem I (PSI).

$$A_j = \frac{(C_c - \Gamma_*)J}{4C_c + 8\Gamma_*} - R_d \quad (3)$$

If adenosine triphosphate (ATP) supply is limited under 100% linear e-transport conditions, factors 4 and 8 in Eq. (3) are replaced by 4.5 and 10.5, respectively [102].

The C4 photosynthetic biochemical model [103] was developed based on the earlier researches [104, 105]. This model fundamentally adapts the core equations of the FvCB model to characterize the metabolic coordination between mesophyll and bundle sheath cells. The pivotal equations characterizing this process are presented as

$$A_c = \frac{(C_s - \gamma_*O_s)V_{\text{cmax}}}{C_c + K_c(1 + O_s/K_o)} - R_d \quad (4)$$

where γ_* is half of the reciprocal of rubisco specificity, C_s and O_s are the CO₂ and O₂ partial pressures in the bundle sheath, respectively.

$$A_j = \frac{(C_s - \gamma_*O_s)(1-x)J_t}{3(C_s + 7/3\gamma_*O_s)} - R_d \quad (5)$$

where J_t is the whole-chain linear e-transport in the mesophyll and bundle sheath chloroplasts, and $(1-x)$ is the partitioning fraction in the bundle sheath chloroplasts.

Since its original publication, the FvCB model has undergone significant evolutionary refinements. A notable limitation of the standard FvCB framework is its inability to account for the photosynthetic insensitivity to fluctuating CO₂ and O₂ concentrations observed under specific environmental conditions. This lack of response is attributed to the rate-limiting step of triose phosphate utilization (TPU) in the synthesis of starch or sucrose, which imposes a third metabolic bottleneck known as TPU limitation [106]. Intensive research into the CO₂ diffusion pathway has further revealed that carbon assimilation is constrained not only by biochemistry but also by stomata and mesophyll conductance [10, 11]. Advanced models, such as eLeaf, have integrated 3D CO₂ reaction-diffusion modules to quantitatively assess how internal leaf anatomy modulates photosynthetic capacity [107]. These studies highlight the critical role of leaf structural plasticity as a target for enhancing crop photosynthesis. Furthermore, to facilitate simulations across diverse thermal regimes, a multitude of temperature-response functions have been developed [108–110]. Through iterative refinements by the scientific community, the FvCB model has emerged as a highly successful tool, demonstrating exceptional reliability in fitting response curves and predicting leaf photosynthetic performance across most scenarios.

Beyond its foundational role in analyzing gas-exchange data [111, 112], the FvCB model has significantly advanced our understanding of electron transport regulation [113] and photosynthetic bottlenecks [114, 115]. By coupling this biochemical framework with stomatal control models [116, 117], researchers have successfully characterized the intricate balance between carbon gain and water loss [118]. Moreover, the model serves as a vital bridge for scaling physiological traits from the chloroplast to the canopy level [119]. This capability is particularly relevant for modern crop improvement, as it enables the quantitative evaluation of how metabolic engineering targets translate into tangible gains in biomass and yield [28, 120].

IV. NEW MISSIONS FOR CANOPY AND LEAF PHOTOSYNTHESIS MODELS

In modern crop models, RUE-based method can result in a 17% overestimation of carbon assimilation [121]. Most importantly, the “black-box” nature of these models precludes their use in identifying molecular targets for crop improvement [26]. To effectively support modern breeding, canopy photosynthesis module in crop models must evolve beyond empirical relationships to incorporate a mechanistic biochemical basis, effectively connecting molecular to whole-canopy productivity.

The engineering of the photosynthesis processes by biotechnologists must be supported by robust biochemical models. While advancements in biotechnology have transitioned traditional breeding toward molecular design breeding, significantly enhancing precision and accelerating cycles, breeders still struggle to identify multi-level physiological traits that effectively guide canopy photosynthesis improvements. Comprehensive canopy photosynthesis models empower researchers to identify diverse potential targets by scenario simulation across multiple scales. A prime example is the acceleration of recovery from photoprotection (non-photochemical quenching), which has been shown to substantially bolster both carbon assimilation and final yield [17]. Furthermore, optimizing leaf chlorophyll content via the reduction of light-harvesting antenna size not only maintains canopy photosynthesis by improving light penetration but also significantly enhances nitrogen use efficiency [13]. *In silico* prediction of crop performance is thus an essential prerequisite before physiologists initiate resource-intensive breeding programs [26, 122]. As a versatile tool for environmental impact assessment, multi-scale canopy modeling allows researchers to quantify the outcomes of specific metabolic or structural modifications [26, 123]. Given the prohibitive costs and logistical limitations of the free-air CO₂ enrichment (FACE) systems, the CoffeePlant3D model provides detailed insights into how coffee trees adapt both structurally and physiologically to carbon-rich atmospheres [124]. These 3D frameworks are particularly adept at characterizing the synergy between irradiance and gas exchange, offering a robust platform for climate-resilient crop design [125].

While existing models demonstrate exceptional performance at either the biochemical or canopy scale, empirical efforts to establish connectivity across these organizational levels remain limited (Fig. 1). To address this, a cross-scale modeling paradigm was pioneered [94], which successfully elevated FvCB model to the canopy scale by integrating the sun-shade leaf approach. Subsequent applications of this

framework have enabled the quantification of potential yield gains from transitioning rice toward a C4 photosynthetic pathway [126], as well as the exploration of how utilizing natural genetic variation can enhance canopy productivity [127]. Furthermore, researchers have evaluated the biochemical bottlenecks limiting light-to-chemical energy conversion efficiency [128], and the impacts of improving photosynthetic properties on crop yield in Europe [120]. Utilizing this method, the revised APSIM model has successfully quantified the impacts of enhancing photosynthesis on crop yield across diverse environments in Australia [28, 129]. Similarly, applications of the revised GreenLab model identified maximal linear electron transport rate and light distribution are the primary limitations in high-density maize production systems [130].

Within this integrative process, coupling biochemical models with the spatiotemporal dynamics of crop development is imperative; without this linkage, simulations are restricted to transient daily predictions rather than seasonal trajectories [119]. Throughout the phenological cycle, foliage undergoes a transition from expansion to senescence, characterized by intricate feedbacks between photosynthetic capacity and internal physiological status [126]. Achieving a molecular-to-canopy modeling pipeline is now a critical priority for accelerating crop improvement in an era of global food insecurity.

V. CONCLUSION

Connecting leaf and canopy photosynthesis models has the capacity to link genetic manipulation of photosynthesis to crop yield. The prerequisite is to ensure that the models at each scale are reliable and effectively linked together. Scaling up the biochemical photosynthesis models to canopy level allows clearer links to the biochemical processes of photosynthesis, based on which consequences of photosynthetic manipulations can be reliably estimated. This can be further improved by the advancing understanding of photosynthetic

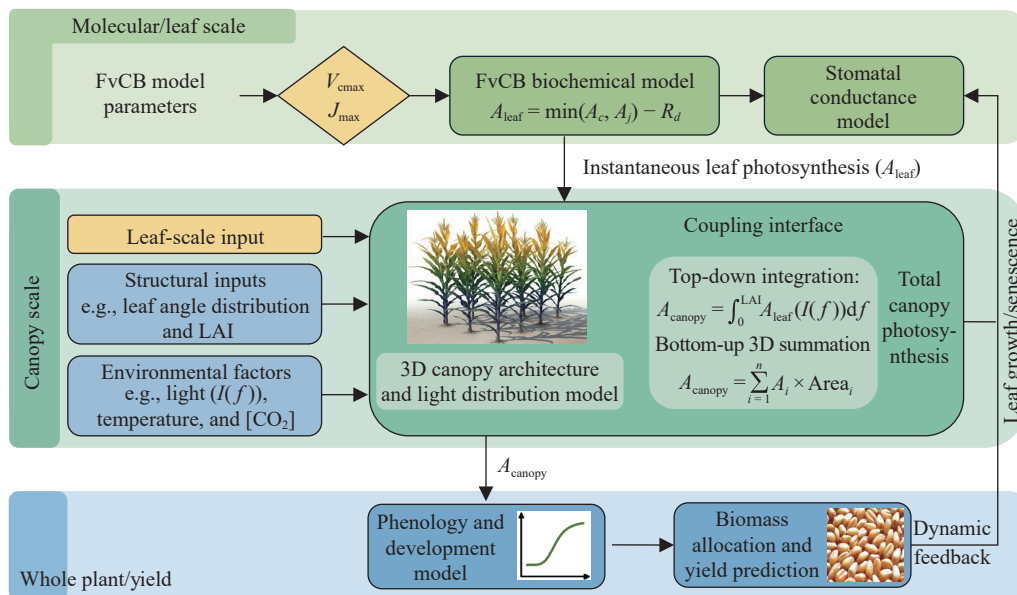


Fig. 1 Multiscale modeling framework linking biochemical to canopy scale.

responses to environmental and physiological attributes and further modeling efforts. Development of the cross-scale modeling framework using the gene-to-phenotype modeling approach can potentially accelerate progress in improving crop resource capture efficiency to support crop improvement through genetic manipulation of photosynthesis.

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