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Title:

A leaf-level biochemical model simulating the introduction of C₂ and C₄ photosynthesis in C₃ rice: gains, losses and metabolite fluxes

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2 **A leaf-level biochemical model simulating the introduction of C₂ and** 3 **C₄ photosynthesis in C₃ rice: gains, losses and metabolite fluxes**

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9 **Summary:**

- 10 • This work aims at developing an adequate theoretical basis for comparing assimilation of the
11 ancestral C₃ pathway with CO₂ concentrating mechanisms (CCM) that have evolved to reduce
12 photorespiratory yield losses.
- 13 • We present a novel model for C₃, C₂, C₂+C₄ and C₄ photosynthesis simulating assimilatory
14 metabolism, energetics, and metabolite traffic at the leaf-level. It integrates a mechanistic
15 description of light reactions to simulate ATP and NADPH production, and a variable
16 engagement of cyclic electron flow. The analytical solutions are compact and thus suitable for
17 larger scale simulations. Inputs were derived with a comprehensive gas exchange experiment.
- 18 • We show trade-offs in the operation of C₄ that are in line with ecophysiological data. C₄ has
19 the potential to increase assimilation over C₃ at high temperatures and light intensities, but this
20 benefit is reversed under low temperatures and light.
- 21 • We apply the model to simulating the introduction of progressively complex levels of CCM
22 into C₃ rice, which feeds more than 3.5 billion people. Increasing assimilation will require
23 considerable modifications such as expressing the NDH complex and upregulating cyclic
24 electron flow, enlarging the bundle sheath, and expressing suitable transporters to allow
25 adequate metabolite traffic. The simpler C₂ rice may be a desirable alternative.

26 **Keywords**

27 Stomata, enzyme, light limitation, C₂ shuttle, C₃-C₄ intermediate, photorespiration, bio-
28 engineering, assimilation.

29 **Running title**

30 Simulating biochemical carbon concentrating mechanisms

31 Introduction

32 Carbon concentrating mechanisms (CCM; acronyms are listed in Table 1) are co-ordinated suites
33 of structural and biochemical modifications to ancestral C₃ photosynthesis. CCMs evolved to reduce
34 the magnitude of photorespiration, a complex process resulting in the release of previously fixed CO₂,
35 which incurs substantial energy costs to recycle by-products (Meyer & Griffiths, 2013). In plants,
36 CCMs have the form of biochemical cycles that increase the CO₂/O₂ ratio at the Rubisco catalytic
37 site, and are of two types: the 'C₂ shuttle' and the C₄ cycle. To operate a CCM, the photosynthetic
38 parenchyma is often differentiated into two cell types, although single-celled systems do exist (King
39 *et al.*, 2012): an external layer of mesophyll (M) and an internal layer of bundle sheath (BS) encircling
40 the vasculature (Lundgren *et al.*, 2014). The C₂ shuttle consists of the compartmentation of glycine
41 decarboxylase (GDC) activity in the BS, delivering CO₂ around Rubisco in the BS, using the
42 photorespiratory glycine produced in the M (Keerberg *et al.*, 2014). The C₄ cycle represents a further
43 sophistication involving an energy dependent carboxylation-decarboxylation cycle. CO₂ is initially
44 fixed into a four-carbon (C₄) organic acid (OAA) in the M by phosphoenolpyruvate carboxylase
45 (PEPC), which after reduction (or transamination) diffuses to the BS where it is decarboxylated. If, on
46 the one hand, the C₄ cycle lowers the photorespiratory ATP demand, on the other it requires a
47 considerable amount of ATP (2 ATP per CO₂ pumped, for the NADP-ME subtype) for the
48 regeneration of phosphoenolpyruvate (Kanai & Edwards, 1999; Evans *et al.*, 2007; Bellasio, 2017).
49 In 'C₂+C₄' species (Bellasio, 2017) the degree of PEPC engagement, and the extent of Rubisco
50 compartmentation to the BS are intermediate and are species dependent (Monson & Moore, 1989). In
51 C₄ species, PEPC is fully engaged and CO₂ accumulates in the BS at concentrations that are 10–
52 20-fold greater than ambient, thereby saturating a fully compartmentalised Rubisco in the BS (von
53 Caemmerer & Furbank, 2003). The biochemical functions of the M and BS need to be separated by
54 a suitable distance (Jurić *et al.*, 2017). Across this space large fluxes of metabolites need to be
55 exchanged, both through plasmodesmata (Osmond & Smith, 1976; Danila *et al.*, 2018), and through
56 a suite of chloroplast membrane transporters (Weber & von Caemmerer, 2010; Gowik *et al.*, 2011;
57 Schlüter *et al.*, 2016).

58 Quantifying the potential gains from operating a CCM has challenged physiologists for the last
59 50 years. Simple approaches have compared C₃ and C₄ plants, but the evolutionary traits of
60 unrelated species can differ substantially, preventing the isolation of the effects of CCMs [reviewed
61 in Snaydon (1991) and Christin and Osborne (2014)]. For instance, in a large comparative
62 experiment Atkinson *et al.* (2016) found C₃ and C₄ grasses mainly differed in terms of leaf mass per
63 area, rather than net assimilation rate per unit leaf area, but Taylor *et al.* (2010) reported that a more
64 limited set of C₄ grasses had a 45 % higher assimilation rate than C₃ grasses. The comparison is
65 further complicated by the co-occurrence of acclimatory traits: Schmitt and Edwards (1981)
66 reported that the effect of short and long term temperature acclimation was greater than any

67 difference in assimilation rate between maize and rice. Even in targeted comparisons between rice
68 and the sympatric weed *Echinochloa glabrescens* or crops such as maize, results were inconclusive
69 (Sheehy, 2007; Covshoff *et al.*, 2016). To quantify the benefit of operating a CCM it is therefore
70 critical to compare two plants in which all traits, other than the strength of the CCM, are equal.

71 For this hypothetical analysis, mathematical models are in principle the ideal tool. Heckmann *et*
72 *al.* (2013) found a smooth monotonic increase in assimilation for increasing levels of C₄ expression
73 in a C₃ background. This finding was directly dependent on the assumption of unlimited ATP, and
74 contrasts with the observation that C₄ plants are favoured only under high temperatures and light
75 intensities (Monteith, 1978; Pearcy & Ehleringer, 1984). Wu *et al.* (2017) compared predictions of
76 C₃ and C₄ models, but these were parameterised separately by curve fitting on representative C₃ and
77 C₄ crops, thereby replicating the unwanted coexistence of multiple traits present in nature within the
78 models. The light-limited model developed by von Caemmerer (2000) assumed a fixed
79 stoichiometric conversion between electron transport and ATP production and is unsuitable for
80 testing different levels of C₄ engagement because the C₄ cycle requires an increased ratio of ATP to
81 NADPH, which C₄ plants obtain by upregulating cyclic electron flow, CEF (Ishikawa *et al.*, 2016).
82 Recently Yin and Struik (2017) overcame some of these shortcomings, but biochemical processes
83 were relatively schematic, and as a result, metabolite exchange requirements have not been
84 quantified.

85 The aims of this work were three-fold. Firstly, to develop the theoretical underpinnings of the
86 introduction of CCMs into C₃ crops at the leaf level; secondly, quantify the possible benefits and
87 trade-offs of CCMs if they were to be made operational in rice; and, finally, estimate realistic
88 fluxes to help define targets for expression of enzymes and transporters. Light-limited formulations
89 working under the assumption of limiting ATP or NADPH, as well as enzyme-limited
90 formulations, all valid for any photosynthetic type, are developed here. These are integrated with a
91 mechanistic description of photosynthetic light reactions, and with a biochemical and
92 hydromechanical model of stomatal behaviour. A gas-exchange experiment was used to inform the
93 model. The results predict that introducing CCMs in C₃ metabolism under the current ambient CO₂
94 concentration would increase assimilation under full light, but the benefit would be reversed at low
95 light intensity (*PPFD*). For C₄ photosynthesis, achieving this potential will require an appropriate
96 electron transport chain, allowing adequate metabolite traffic, and enlarging the BS to house the
97 biochemical and light harvesting machinery.

99 **Material and Methods**

100 *Overview of the modelling approach*

101 The modelling scheme is depicted in Figure 1 to highlight key inputs and outputs, This model
102 was newly derived to allow a seamless transition between all photosynthetic types except CAM,
103 and joins together an electron transport submodel, a biochemical submodel, a stoichiometric
104 submodel (see schematic in Figure S1), and a stomatal submodel. The photosynthetic type is
105 defined by setting the strength of the C₄ cycle [as PEP carboxylation rate ($V_{P(J)}$) in the light-limited
106 model and maximum rate of PEPC, ($V_{P\text{MAX}}$) in the enzyme-limited submodel] together with the
107 location of GDC (χ_{GDC}). The electron transport submodel (Note S1, Figure S2) calculates the flux of
108 ATP and NADPH (J_{ATP} and J_{NADPH}) made available under a given *PPFD*. Here, the limitations of
109 previous modelling approaches using a fixed stoichiometry of the electron transport chain (see
110 *Introduction*) were resolved by allowing the ratio of ATP/NADPH production to be adjusted
111 through mechanisms that were found to be critical in C₄ plants. These are the regulation of the rate
112 of cyclic electron flow (CEF) through the parameter f_{Cyc} , and inducing the NAD(P)H
113 Dehydrogenase-like (NDH) complex (Ivanov *et al.*, 2005; Friso *et al.*, 2010; Munekage *et al.*,
114 2010) which is characteristic of C₄, and not used by C₃ plants, operating mainly the PGR5 / PGRL1
115 pathway (Yamori & Shikanai, 2016) by varying f_{NDH} (the fraction of CEF passing through the NDH
116 complex). The reducing power requirements of nitrogen reduction are implicitly accounted for here
117 as pseudocyclic electron flow (lumped with the water-water cycle, and adjusted through $f_{\text{Pseudocyc}}$),
118 in line with Yin and Struik (2012).

119 The biochemical submodel has different formulations depending on the limitation, sharing
120 common underpinnings (Note S2). There is a formulation for limitation by Rubisco or PEPC
121 carboxylating capacity (commonly referred to as enzyme limitation, Note S3) and two formulations
122 for light-limited photosynthesis, derived under limiting ATP (Note S4) or NADPH (Note S5).
123 Equations for triose phosphate limited photosynthesis (Busch *et al.*, 2018) were omitted for
124 simplicity as they are relevant under low O₂ or high CO₂ concentrations, or low temperatures
125 (Busch & Sage, 2017), while crops like rice – fertilised and irrigated – generally experience mainly
126 light limitations (Yin & Struik, 2015). Similarly, limitations imposed by the diffusion of
127 metabolites (Retta *et al.*, 2016) were neglected for simplicity, justified by a recent study addressing
128 the introduction of a weak C₄ cycle in C₃ photosynthesis using a reaction diffusion model that found
129 that any reduction of *A* due to the effect of diffusion processes was limited (Wang *et al.*, 2017). The
130 ATP and NADH produced during respiration were neglected because they are likely to be
131 consumed by basal metabolism, while NADH imbalances are likely to be dissipated by
132 mitochondrial alternative oxidases (Buckley & Adams, 2011).

133 Using dummy values (initial values for a converging iteration) for the CO₂ concentration at the
134 M carboxylating sites (C_M) the light-limited submodel calculates two distinct sets of outputs, under

135 NADPH and ATP limitations. Of those, that resulting in the minimum V_C is taken as output of the
136 light-limited model. Similarly, starting from C_M , the enzyme-limited submodel calculates a full set
137 of outputs using the kinetic characteristics of Rubisco and PEPC as inputs.

138 Outputs of light-limited and enzyme-limited submodels are joined using a smoothing function
139 to give a continuous output (Note S6), as well as used to calculate τ , a quantity related to the ATP
140 concentration in the M and the BS that acts as the biochemical driver of stomatal response (Note
141 S7). This was included solely to realistically simulate stomatal conductance in a C_3 to C_4
142 continuum, but we make no claim about whether τ offers a faithful mechanistic description of
143 stomatal behaviour. Hydro-mechanical forcing links guard cell responses to the water status and
144 turgor of the leaf, which relate to soil water status and plant hydraulic conductance. The influence
145 of biochemical factors relative to hydro-mechanical forcing is determined by the parameter β , while
146 stomatal morphology is described by χ_S . The output of the stomatal submodel is stomatal
147 conductance, (g_S) that, together with mesophyll conductance g_M , is used to calculate C_M , which is
148 iterated. Temperature dependence is simulated with empirical functions (Note S8, Table S2). For
149 each combination of inputs, the locality of Rubisco between BS and M (χ_{Rubisco}) together with the
150 rate of flow through CEF (f_{Cyc}) were fitted to maximise A . This resulted in light reactions generating
151 exactly the ATP and NADPH which was consumed by dark reactions, while the ATP-limited
152 model and the NADPH-limited models converged to output the same level of A . The outputs of
153 these submodels (V_{OBS} , V_{CBS} , V_{OM} , and V_{CM}) were inputted to a generalised stoichiometric model of
154 assimilation (Bellasio, 2017), used to calculate reaction rates, and fluxes across the BS and M
155 interface (Figure S1). Here, three additional inputs partition key processes between the BS and M:
156 f_{PR} , for phosphoglycerate reduction; f_{CS} , for carbohydrate synthesis; f_{PPDK} , for pyruvate phosphate
157 dikinase (Table 1). Model parameterisation and sensitivity are described in Notes S9 (coefficients
158 are in Table S3) and S10, respectively.

159 *Plants, gas exchange, and fluorometry*

160 Plants of *Oryza sativa* subsp. *indica*, modern, high-yielding variety Takanari (Taylaran *et al.*,
161 2009) were germinated and grown in 1.5 L pots filled with Martins potting mix (80% composted
162 bark, 10% coir, 10% sand, complete fertiliser), in acrylate greenhouses located in Canberra (35°S,
163 149°E) under natural illumination in April – May 2018. Pots were partially submerged for a third of
164 the depth in polypropylene tubs and watered weekly for six weeks. Gas exchange and fluorescence
165 were measured on a fully expanded leaf with a setup similar to Bellasio and Griffiths (2014b).
166 Briefly, a portable gas exchange system (LI6400XT, Li-Cor, Lincoln, USA) was modified to
167 operate at low CO_2 concentrations (see licor.com) and fitted with a 6400-06 PAM2000 adapter,
168 holding a fibre probe in the upper leaf cuvette distant enough to avoid shading. Light was provided
169 by a bespoke red-blue light source, positioned to illuminate uniformly the leaf. Light intensity was
170 measured through an in-chamber Gallium arsenide photodiode, calibrated using a Li-250 light

171 sensor (Li–Cor). Neoprene gaskets were used on both sides of the cuvette. A mixture of 2 % O₂ was
172 prepared by mixing ambient air and N₂ with a bespoke gas mixing unit (kindly assembled by Suan
173 Chin Wong). This mix or ambient air was CO₂–scrubbed with soda lime and humidified to a dew
174 point of 15–17 °C upstream of the inlet to maintain water vapour pressure deficit around 1 kPa. CO₂
175 was added from a cylinder (Isi, Vienna, Austria), using the CO₂ injection unit of the LI6400XT.
176 PSII yield was measured with a Dual PAM–F (Heinz Walz GmbH, Effeltrich, Germany). Pulse
177 intensity was adjusted to be between 10,000 and 12,000 μmol m⁻² s⁻¹ thereby exceeding the
178 requirements of between 6,000 and 8,000 μmol m⁻² s⁻¹, depending on CO₂ and *PPFD* levels, to
179 saturate the fluorescence signal. Mass flow leaks (Boesgaard *et al.*, 2013) were monitored with a
180 gas flow meter as detailed in Bellasio, C. *et al.* (2016), and sealed with a tiny ridge of atoxic
181 gelatine laid between the gaskets and the leaf. Four photosynthetic response curves were measured
182 at 25 °C on *n*=4 plants as detailed in Bellasio, C. *et al.* (2016). *A/C_i* curves were measured under a
183 *PPFD* of 1200 μmol m⁻² s⁻¹, light curves were measured under a *C_a* of 420 μmol mol⁻¹. Flow rate
184 was 490 μmol s⁻¹; CO₂ diffusion through the gaskets was compensated by lengthening the tubing of
185 the LI6400XT reference gas.

186 **Results**

187 *Gas exchange*

188 The operational conditions of rice plants were characterised by a comprehensive gas exchange
189 experiment, which combined measurements under ambient and low O₂. Primary, diffusion leak–
190 corrected data appear as symbols in Figure 2, PSII yield is shown in Figure S3. Overall, rice
191 displayed typical C₃ responses. Under high *PPFD* (Figure 2A), *A* was lower under ambient O₂
192 (closed symbols) than under low O₂ (open symbols) because of photorespiration. The quantum yield
193 for assimilation (the initial slope of the curves), was higher under low O₂ (0.0397±0.0002 and
194 0.0512±0.0023 under ambient and low O₂, respectively). Under low *C_i* (Figure 2B), *A* was higher
195 under low O₂ than under ambient O₂ because of O₂ competitive inhibition of Rubisco. Assimilation
196 saturated at relatively lower *C_i* under low O₂ (open symbols) than under ambient O₂. The stomatal
197 conductance (*g_s*) measured in *A/PPFD* curves (Figure 2C) increased monotonically with *PPFD*
198 showing a saturating response similar to that of the *A/PPFD* curve. Under varying external CO₂
199 concentration (*C_a*), *g_s* decreased non–linearly with slope depending on the O₂ level. Rice had a
200 slightly higher *in vivo* *S_{C/O}* (Table 1) than that found *in vitro* (Hermida-Carrera *et al.*, 2016) perhaps
201 for the tight association between mitochondria and chloroplasts that evolved to maximise
202 photorespiratory CO₂ recapture (Sage & Sage, 2009; Hatakeyama & Ueno, 2016). Under a *PPFD*
203 of 500 μmol m⁻² s⁻¹, rice operated at a relatively low *V_O/V_C* of circa 0.3 [Figure S4, compare with
204 Bellasio *et al.* (2014)].

205 *Simulating assimilation and stomatal conductance of native C₃ rice*

206 A/C_i and $A/PPFD$ curves responses for rice were simulated in the same conditions used for gas
207 exchange measurements. The model predicted with accuracy $A/PPFD$ (Figure 2A) and A/C_i curves
208 (Figure 2B) measured under ambient O_2 , but overestimated $A/PPFD$ and A/C_i curves under low O_2
209 and high C_a . We attribute this to triose phosphate limitation, and to the feedbacks regulating the
210 electron transport chain through the quenching of $Y(II)$ under low O_2 (Figure S3) which we have
211 addressed in Bellasio (2018) but not considered in this model, for simplicity. The simulated
212 stomatal behaviour captures very well the shape of the stomatal response, in both $A/PPFD$ and A/C_i
213 curves and at both O_2 levels.

214 *Simulating gas exchange of C_2 , C_2+C_4 and C_4 rice*

215 Here, simulations were intended to capture hypothetical best-case scenario, assuming unlimited
216 phenotypic plasticity whereby Rubisco is optimally distributed and electron transport processes
217 fully accommodate CEF and NDH levels. Conditions and fitting routines were the same as used for
218 the C_3 simulations. The C_2 shuttle and progressive levels of C_4 activity were introduced in native
219 rice by manipulating the activity of PEPC (through the inputs $V_{P_{MAX}}$ and $V_{P(J)}$), the locality of GDC
220 (ξ_{GDC}), the engagement of the NDH pathway of electron transport (f_{NDH}) and the BS apportioning of
221 light respiration (f_{RLIGHT} , see Table 1 for full details). The levels of the fitted inputs $\chi_{Rubisco}$ and f_{Cyc}
222 are shown in Figure S5. These are relevant for bioengineering as they indicate the required physical
223 distribution of Rubisco, and the necessary adjustments to the electron transport chain. $A/PPFD$
224 curves (Figure 3A) simulated at a C_a of $400 \mu\text{mol mol}^{-1}$ intersect around a $PPFD$ of $300 \mu\text{mol m}^{-2} \text{s}^{-1}$.
225 Under lower $PPFD$ s C_2 A was the highest and C_4 was the lowest. Under higher $PPFD$ s A
226 increased proportionally with the level of CCM engagement and was $\sim 22\%$ higher for C_4 than C_3 at
227 a $PPFD$ of $1500 \mu\text{mol m}^{-2} \text{s}^{-1}$. The analysis of A/C_i curves (Figure 3B) revealed expected
228 differences in predicted gas change characteristics between photosynthetic types, with A at C_a lower
229 than $\sim 550 \mu\text{mol mol}^{-1}$ being progressively higher for plants operating CCMs at increasing
230 engagement. But the operation of a CCM necessarily sacrifices A under higher C_a . There were
231 striking differences in stomatal conductance, which was around 40% less in C_4 than in C_3 under a
232 $PPFD$ of $1500 \mu\text{mol m}^{-2} \text{s}^{-1}$ and a C_a of $400 \mu\text{mol mol}^{-1}$ (Figure 3C), indicating that the same level
233 of A was achieved with lower transpiration and higher water use efficiency, in line with differences
234 between extant C_3 and C_4 species (Bellasio *et al.*, 2018; Quirk *et al.*, 2018) although in the field
235 there is some negative feedback on the effect on WUE because of temperature changes. The same
236 differences were maintained in the simulated A/C_i curves (Figure 3D). Notably these differences in
237 g_s resulted solely from biochemical differences between photosynthetic types (sensed by the
238 quantity τ) while all other parameters were maintained at C_3 levels. The operation of the CCMs
239 resulted in an increase in the CO_2 concentration in the BS (Figure 3E and 3F) and in the consequent
240 reduction of the ratio between Rubisco oxygenation and carboxylation (Figure 3G and 3H). The
241 output fraction of BS Rubisco carboxylation V_{CBS}/V_C , which depends both on C_{BS} and on $\chi_{Rubisco}$, is

242 shown in Figure 3I and 3J. V_{CBS}/V_C was relatively invariant with $PPFD$ in all photosynthetic types
243 except C_4 , where it slightly decreased below $500 \mu\text{mol m}^{-2} \text{s}^{-1}$ (Figure 3I). In A/C_i curves V_{CBS}/V_C
244 increased at low C_a for C_2 and C_2+C_4 types and decreased at high C_a for the C_4 type. Leakiness (the
245 rate of CO_2 retrodiffusion from the BS relative to PEP carboxylation rate), of relevance for isotopic
246 studies, (Cernusak *et al.*, 2013; Bellasio & Griffiths, 2014b) is plotted in Figure S6. To isolate any
247 effect of CO_2 diffusion through the mesophyll and stomata, these simulations were repeated using
248 C_M as input, and are shown in Figure S7.

249 *Assimilatory gain/loss of C_2 , C_2+C_4 and C_4 rice at different temperatures, C_a , and $PPFD$*

250 This set of simulations explored gains and losses of operating different types of photosynthesis,
251 as compared to C_3 . Three scenarios were simulated: one of unlimited plasticity of the electron
252 transport chain and two in which some elements of the electron transport chain remain in a C_3
253 configuration. In the best case scenario electron transport processes fully accommodate the ATP
254 demand of different types of CCM through the optimisation of the levels of CEF (f_{Cyc}) and by
255 allowing expression of the NDH complex in C_2+C_4 and C_4 types ($f_{NDH}>0$). Figure 4 shows that
256 operating C_2 was beneficial at all temperatures and $PPFD$ s, but gains were generally lower than
257 10% (Figure 4B), as compared to C_3 (Figure 4A). Operating C_2+C_4 was slightly counterproductive
258 below a $PPFD$ of $450 \mu\text{mol m}^{-2} \text{s}^{-1}$ and a temperature of 40°C but allowed substantial gains above
259 (Figure 4C). The range in which operating C_4 photosynthesis did not confer net benefits was cutting
260 diagonally below a temperature of 40°C and a $PPFD$ of $500 \mu\text{mol m}^{-2} \text{s}^{-1}$ (Figure 4D). The possible
261 gains and losses were much more pronounced for C_4 than for C_2 and C_2+C_4 types. In the operation
262 of the C_4 cycle most of the energy saved by suppressing photorespiration is consumed by the
263 regeneration of PEP; the resulting balance depends on their relative flux, and can be quantified
264 through the quantum efficiency of assimilation $Y(\text{CO}_2)$, shown on incident light basis in Figure S8.
265 $Y(\text{CO}_2)$ was very similar for C_3 and C_2 types. C_2+C_4 and C_4 had higher $Y(\text{CO}_2)$ than C_3 at high
266 $PPFD$ s, but lower at low $PPFD$ s. Overall, $Y(\text{CO}_2)$ was slightly lower than our previous
267 measurements in tobacco and maize (Bellasio, C. *et al.*, 2016; Bellasio, Chandra *et al.*, 2016),
268 which we attribute to slightly lower $Y(II)_{LL}$ and s (Table 1).

269 We then compared CCM types to C_3 assimilation in the temperature and C_a space, under a
270 moderate $PPFD$ of $700 \mu\text{mol m}^{-2} \text{s}^{-1}$, meant to capture illumination of an ordinary erect leaf of a
271 modern cultivar in the upper level of the canopy, in the same optimistic scenario of variable CEF
272 and engaged NDH (Figure 4E). C_2 assimilation was beneficial at all temperatures and C_a (Figure
273 4F). Gains were greater than 10% in a relatively broad set of conditions including under ambient C_a
274 at high temperatures. The C_4 and C_2+C_4 types were disadvantageous above a C_a of around 450
275 $\mu\text{mol mol}^{-1}$ and below 40°C – a broader range than under higher $PPFD$ (Figure 3B). The C_4 and
276 C_2+C_4 types were progressively more advantageous at higher temperature and low C_a .

277 Similar simulations were carried out to represent a less optimistic scenario whereby the activity
278 of the NDH complex remained at C₃ levels ($f_{\text{NDH}}=0$) for all photosynthetic types (Figure 5, top
279 row). The marginal gains were maintained for the C₂ type (Figure 5A); however, C₂+C₄ and C₄
280 types were counterproductive in a broader range of *PPFDs* roughly cutting below a *PPFD* of 700
281 $\mu\text{mol m}^{-2} \text{s}^{-1}$ for the C₂+C₄ type and 900 $\mu\text{mol m}^{-2} \text{s}^{-1}$ for the C₄ type (Figure 5B and 5C).

282 In a pessimistic scenario, in addition to the incapacity to express sufficient NDH complex
283 ($f_{\text{NDH}}=0$), CCM types were unable to modify the flux through CEF, which remained capped at C₃
284 levels (Figure 5, bottom row). Here, the marginal gains were maintained for C₂ photosynthesis
285 (Figure 5D); however, the C₂+C₄ type was counterproductive below a *PPFD* of 1000 $\mu\text{mol m}^{-2} \text{s}^{-1}$,
286 while the C₄ type was counterproductive at all *PPFDs* below a temperature of 30°C (Figure 5E and
287 5F). Severe losses in excess of 40% were predicted for the C₄ type at ordinary temperatures and
288 moderate to low *PPFDs*.

289 *Metabolite transport*

290 Two further sets of simulations estimated the metabolite fluxes between the M and the BS by
291 manipulating the level of C₄ engagement through increasing levels of V_P (Figure 6) so as to
292 represent the full C₂+C₄ continuum from C₂ (left of each panel) to C₄ (right of each panel). In a first
293 scenario (Figure 6A), the level of ATP demand in the BS was minimised. In these conditions,
294 phosphoglycerate is not reduced in the BS but diffuses to the M and is reduced therein to
295 dihydroxyacetone phosphate, DHAP. A minimal part of DHAP is used by carbohydrate synthesis,
296 but the majority diffuses back to BS to replenish the sugar phosphates pool. This drives the
297 metabolite exchange between the M and the BS to a maximum. In addition, because
298 phosphoglycerate reduction is the main NADPH sink in the BS, when ATP demand in the BS is
299 minimal, the NADPH demand in the BS is also minimal. This requires by-passing the malate
300 dehydrogenase in the M, and, to maintain the efficiency of the CCM despite the inability to operate
301 the malate shuttle, the CCM works through alanine and aspartate (Bellasio, 2017). This condition is
302 suboptimal because it requires high concentration gradients of aspartate and alanine when
303 malate and pyruvate do not transport CO₂ (Arrivault *et al.*, 2017). At low levels of C₄ engagement,
304 when V_P was low, glycine and serine were operating the C₂ shuttle. The model predicts that the
305 reducing power generated in the BS by the decarboxylation of glycine, which could not be used by
306 phosphoglycerate reduction because of the insufficient ATP availability, was returned to the M by
307 the malate and pyruvate shuttle in a 'backward' C₄ cycle. As V_P increased, the flux of glycine and
308 strength of the C₂ cycle [which scales with V_P , see details in Bellasio (2017)] was progressively
309 reduced, diminishing the excess NADPH in BS together with the malate and pyruvate fluxes that
310 decrease to zero with V_P . With the increase in V_P , the fitted fraction of Rubisco carboxylation in BS
311 increased linearly, causing the ratio of ATP demand in BS relative to M to increase linearly (Figure
312 6C).

313 An opposite scenario, where fluxes were minimal, was simulated by fitting f_{PR} and f_{CS} to
314 minimise the sum of squared flow rates between BS and M (Figure 6B). In these conditions the
315 increase of phosphoglycerate reduction in the BS drove the ATP demand in the BS to a maximum
316 (Figure 6D). The total fluxes were less than half those of the previous case (54 *versus* 130 $\mu\text{mol m}^{-2}$
317 s^{-1}); the main metabolites to be transported in these conditions were malate and pyruvate, which
318 were the sole compounds to support the CCM while the flux of aminoacids was minimal. Despite
319 the malate and pyruvate shuttle working in full, and exporting reducing power from the M to the
320 BS, the NADPH demand in the BS was high (Figure 6D), requiring substantial linear electron flow
321 in the BS ($\sim 18 \mu\text{mol m}^{-2} \text{s}^{-1}$ of NADPH).

322 Discussion

323 This work set out to study the theoretical underpinnings of the introduction of CCMs into C_3
324 metabolism. A model of enzyme and light-limited assimilation was newly derived to account for
325 the stoichiometry of Bellasio (2017) (Table S1) augmented to include the explicit mechanistic
326 description of the electron transport chain (Bellasio, 2018), and a hydromechanical and biochemical
327 model of stomatal conductance recently shown to work for C_3 and C_4 plants (Bellasio *et al.*, 2017).
328 We shall stress four points distinguishing the importance of this work. Firstly, by including a
329 hydromechanical submodel we provide a means to connect plant assimilatory biochemistry to plant
330 hydraulics, allowing the concurrent investigation of photosynthesis and water use. Secondly, this is
331 the only study comparing C_2 performance with C_3 , C_2+C_4 and C_4 seamlessly within a single model,
332 offering a further improvement over approaches targeted to specific types. Thirdly, this is the only
333 study estimating the metabolite fluxes necessary to operate the different photosynthetic types.
334 Lastly, the model marries biochemically comprehensiveness (it includes all main reactions of the
335 photosynthetic metabolism) with computational speed, required by larger scale modelling. This
336 model is generally applicable, and will be valuable for ecophysiological and evolutionary studies,
337 but we will address evolution at a later stage. Here, we applied the modelling framework to predict
338 assimilation and metabolite fluxes in a three dimensional environmental landscape ($t \times C_a \times PPF$)
339 using parameters derived for rice. Next, we make some general considerations on the introduction
340 of a CCM in C_3 metabolism, and we elaborate on the special case of rice.

341 There is a pervasive belief that the introduction of C_4 photosynthesis into C_3 plants will
342 unconditionally increase assimilation, supported by models based on the assumption that ATP and
343 NADPH are unlimited (Heckmann *et al.*, 2013). However, decades of comparison between C_4 and
344 C_3 plants have shown that C_3 plants may be advantaged in a range of conditions [e.g. (Ehleringer *et al.*,
345 1997; Ghannoum *et al.*, 2000; Christin & Osborne, 2014)]. We showed that, when energy
346 budgets were accounted for, C_4 photosynthesis becomes unfavourable at high CO_2 concentrations,
347 low $PPFD$ and low temperatures, and therefore provide a novel theoretical framework to explain
348 such experimental observations.

349 *Bundle sheath permeability mediates trade-offs imposed by light intensity*

350 Modern crops like rice have typically a LAI (leaf area per ground area) of 5–6, meaning that the
351 majority of leaves are shaded and, importantly, the overall performance of C₄ types will
352 compromise full-light advantages and shade disadvantages. The key parameter governing
353 photosynthetic losses under low *PPFD* in C₄ photosynthesis is BS conductance, g_{BS} (Bellasio &
354 Griffiths, 2014b). g_{BS} controls the flux of CO₂ released in the BS that retrodiffuses to the M, called
355 leakage (Farquhar, 1983). g_{BS} can vary several orders of magnitude in nature and can affect *A*
356 substantially (Kromdijk *et al.*, 2014; Yin & Struik, 2017), in particular at high levels of CCM
357 engagement (Figure S9). Under high temperature, g_{BS} is reported to increase (Yin *et al.*, 2016),
358 while under low *PPFD* V_P decreases, driven by a reduced rate of ATP production (Bellasio &
359 Griffiths, 2014b). In these conditions, leakage reduces C_{BS} , and, in C₄ plants, it dissipates energy
360 through the ATP-dependent regeneration of phosphoenolpyruvate required to re-fix the leaked
361 CO₂, making the CCM counterproductive (Tazoe *et al.*, 2008; Ubierna *et al.*, 2011; Ubierna *et al.*,
362 2013; Bellasio & Griffiths, 2014b; Sun *et al.*, 2014; Pignon *et al.*, 2017). In nature, plants minimise
363 the ratio between leakage and metabolite fluxes by preferentially localising plasmodesmata at the
364 interface between M and BS, while apoplastic diffusion is often reduced by the deposition of a gas-
365 tight suberized cell wall (Sowinski *et al.*, 2008; Sowiński, 2013; Danila *et al.*, 2016; Danila *et al.*,
366 2018). If low g_{BS} may therefore appear desirable (though perhaps difficult to achieve), high
367 symplastic permeability is required to sustain metabolite diffusion [Figure 6, (Weber & von
368 Caemmerer, 2010)], and this dilemma constitutes an efficiency trade-off that is inherent to the C₄
369 CCM – and unavoidable (Bellasio & Griffiths, 2014a). Indeed, to attune leakage to *PPFD* levels,
370 g_{BS} in maize was found to adjust during growth (Bellasio & Griffiths, 2014b) as well as in adult
371 leaves (Bellasio & Griffiths, 2014a).

372 *Future CO₂ levels*

373 Rising anthropogenic atmospheric CO₂ concentrations will favour C₃ assimilation over C₄. Apart
374 from the difficulties in predicting future CO₂ levels – not addressed here – predicting assimilation
375 under changing CO₂ is very difficult. When plants are exposed to a high CO₂ level for a long time
376 they may downregulate the pool of Rubisco and PEPC (Ghannoum *et al.*, 2000; Leakey *et al.*, 2004;
377 Long *et al.*, 2006; Leakey *et al.*, 2012), at the same time, producing fewer stomata (Way *et al.*,
378 2011; Franks *et al.*, 2012)(Quirk, Bellasio and Beerling, *Annals of Botany*, *in press.*). There is a
379 growing body of data gained under controlled conditions [e.g. (Bellasio *et al.*, 2018; Quirk *et al.*,
380 2018)] and in free air experiments [e.g. (Bishop *et al.*, 2015)], yet, responses are species specific
381 and, currently, evidence is not sufficient to generalise acclimation responses of C₄ and C₃ plants. As
382 a result, it is common practice in climate modelling to take assimilatory responses measured under
383 transient changes in CO₂ levels (*A/C_a* curves) as predictive of stable responses of plants grown
384 under different CO₂ levels, that is, no large scale models include representation of the physiological

385 acclimation to future CO₂ level (Rogers *et al.*, 2017). With this principle, using simple interpolation
386 of the best case scenario shown in Figure 4H, at 25 °C, C₄ assimilation would equal C₃ assimilation
387 at a C_a of 465 μmol mol⁻¹, a level that would be exceeded in 2036 according to the A2 scenario of
388 carbon emission mitigation (http://www.ipcc-data.org/observ/ddc_co2.html).

389 *Strategies for engineering a CCM*

390 In the face of global warming, the introduction of CCMs in a C₃ crop such as rice was proposed
391 as a possible strategy to increase yield (Leegood, 2013; Long *et al.*, 2015). An operational C₂
392 shuttle was considered as a first step in bio-engineering, with the final goal of obtaining a fully
393 expressed C₄ type. Of the three biochemical C₄ subtypes (NADP-ME, NAD-ME, PEPCK), the
394 NADP-ME was chosen as the initial target (Kajala *et al.*, 2011), as it is operated by the crops with
395 greatest productivity (Furbank, 2011) and would require introducing a smaller number of enzymes
396 [in M cells carbonic anhydrase, PEPC, malate dehydrogenase, and pyruvate-phosphate dikinase; in
397 BS cells NADP-ME, plus eight transmembrane transporters (Kajala *et al.*, 2011)]. Other subtypes
398 require additional enzymes [aspartate and alanine aminotransferase, PEPCK, NAD-ME (Wang *et al.*
399 *et al.*, 2014), plus up to three transporters (Schlüter *et al.*)] and were not considered here, but see
400 Bellasio (2017). Traditionally, strategies for engineering a CCM have emphasized the manipulation
401 of dark reactions and the associated genetics (Kajala *et al.*, 2011; Leegood, 2013). Here we point to
402 two overlooked factors required for the operation of a CCM, namely anatomy and light reactions.

403 Firstly, leaf anatomy needs to be adjusted depending on the level of C₄ cycle expression.
404 Anatomy and biochemistry of the BS are mutually interdependent (Bellasio & Griffiths, 2014c).
405 The requirement in light harvesting optical cross section depends on the ATP demand, and
406 determines the required BS volume, mediated by the size of the ATP-generating light harvesting
407 machinery, plus the volume of the dark reactions machinery (Bellasio & Lundgren, 2016). Minimal
408 ATP demand in the BS may be desirable as it would require the smallest BS, and therefore require
409 minimum modification of the current rice anatomy, but would lead to the unwanted necessity of
410 high gradients and flux rates, and require the expression of high levels of metabolite transporters
411 (Pick *et al.*, 2011). Aiming at a high ATP demand would have the benefit of requiring the minimum
412 expression of transporters but would require the largest electron transport chain, and therefore a
413 more radical modification of the native C₃ anatomy. Identifying a desired anatomical target requires
414 therefore first to identify a biochemical ideotype. Each of the two extreme solutions shown in
415 Figure 6 would entail limited operational robustness (Pick *et al.*, 2011), as there would not be any
416 freedom to accommodate transient environmental change (Bellasio & Griffiths, 2014c). A ‘robust
417 flexibility’ would be positioned half-way between these two opposite scenarios, for instance where
418 the ATP demand in the BS relative to M is 0.7. The potential ratio of ATP production in the BS
419 relative to M must exceed 0.7 by a considerable safety margin (Bellasio & Lundgren, 2016) to
420 counter changing light conditions (Bellasio & Griffiths, 2014c). To achieve this, the light absorbed

421 in the BS relative to M under white light, must be close to 0.7. Currently, the size and pigmentation
422 of rice BS is insufficient (Bellasio & Lundgren, 2016). A suitable situation was found in maize,
423 which had a BS pigmentation circa twice that of the M, and allocated ~30% of the total leaf section
424 area to the BS (Bellasio & Lundgren, 2016) and should be considered as the target for C₄ rice.
425 Further, reaching the required levels for g_{BS} will require engineering the appropriate density of
426 plasmodesmata (Danila *et al.*, 2016), reducing leakage, and possibly allow for acclimation of g_{BS}
427 during growth (see above). Alternatively, higher efficiency could be reached by operating the C₄
428 cycle only in those parts of the canopy where the *PPFD* is higher than a given threshold, but this
429 seems difficult to achieve also because it is adopted neither in mature nor in developing maize
430 leaves (Wang *et al.*, 2013).

431 Secondly, the operation of a C₄ cycle will require important modifications to the electron
432 transport chains. We showed that when cyclic electron flow, CEF (f_{Cyc}) and the NDH pathway
433 (f_{NDH}) were allowed to vary (Figure 4), the performance of C₂+C₄ and C₄ types was maximal. This
434 optimal scenario reflects the idea that electron transport processes may spontaneously adjust in
435 response to the expression of a CCM, responding to an increase in ATP demand, through flexibility
436 mechanisms inherent in native chloroplasts (Takeuchi *et al.*, 2000). Higher levels of f_{NDH} would
437 benefit C₄ assimilation, but may be physiologically implausible, for example because NDH is very
438 expensive to produce and maintain. It is possible, however, that rice does not have the potential to
439 express adequate level of CEF and NDH components. If f_{NDH} is capped at C₃ levels the performance
440 of C₄ rice will be lower (Figure 5 A–C), and if f_{Cyc} is capped at C₃ levels *A* would be depressed even
441 further (Figure 5 D–F).

442 Considering the complexities and trade-offs of implementing a C₄ cycle, C₂ rice may be a
443 desirable product of bioengineering efforts. Despite the relative operational simplicity, the
444 engagement of a C₂ shuttle always increased assimilation rate, relative to C₃. The assimilation gain
445 was relatively small under ambient C_a , but increased with temperature at low C_a (Figure 4F).
446 Although in water-rich rice paddies plants can maintain stomata open and extreme photorespiratory
447 conditions might not occur at mid-latitudes (where temperatures are milder and the subsp. *japonica*
448 is favoured), they may occur at low-latitudes (where temperatures are higher and the subsp. *indica*
449 is favoured), and, particularly, for dryland rice, which would probably be the crop to benefit most
450 from the introduction of a C₂ CCM. In the simulations, the locality of Rubisco activity, as $\chi_{Rubisco}$,
451 was adjusted continuously at varying C_M always resulting in optimal Rubisco activity. In nature,
452 however, the proportion of Rubisco in the BS may change only on evolutionary timescales and may
453 be plant-specific. Consequently, there may be a trade-off between optimisation for
454 photorespiratory conditions, by compartmentalising more Rubisco to the BS, or for non-
455 photorespiratory conditions by allowing all Rubisco in the M, with easier access to intercellular
456 CO₂. Allocating 10 % of Rubisco in the BS was a good compromise (Figure S5).

457 *From leaf-level to crop*

458 Upscaling these findings to calculate crop yield will be a challenging task. Firstly, it will require
459 modelling of the canopy light environment (Song *et al.*, 2013), possibly including diel light cycles
460 of fully illuminated leaves (Wu *et al.*, 2017) and the transient illumination in shaded leaves (Percy
461 *et al.*, 1997), nitrogen allocation (Buckley *et al.*, 2002; Dewar *et al.*, 2012), the effect of different
462 canopy architectures (Burgess *et al.*, 2017), the response of A and g_s to temperature and humidity
463 (Yin & Struik, 2017). Ideally, the description could consider the potential losses due to suboptimal
464 stomatal aperture (Violet-Chabrand *et al.*, 2016; Bellasio *et al.*, 2017), and the mid-morning
465 depressions of photosynthetic capacity (Horton & Murchie, 2000). The necessity of translating
466 assimilation into grain yield will add further complexities and require a dedicated crop model
467 accounting for root growth, nitrogen uptake, pathogens, as well as the interactions between cultivars
468 and climate (Li *et al.*, 2015; Paleari *et al.*, 2017). There is an urgent need for addressing some of
469 these challenges. This model offers the necessary underpinnings and can be readily used as a
470 submodel for modelling assimilation at higher spatial level.

471 **Conclusion**

472 We developed new ATP-limited, and NADPH-limited submodels of assimilation, as well as a
473 light reaction submodel, coupled with a stomatal submodel. The resulting model connects light
474 harvesting to dark assimilatory biochemistry and hydraulics and is valid for any photosynthetic
475 type. The equations were solved analytically and will be valuable for evolutionary as well as
476 ecophysiological studies, and we encourage their use also for larger scale modelling. The model
477 was calibrated and tested on primary gas exchange and fluorescence data measured on rice. By
478 simulating the introduction of CCMs in C_3 metabolism we showed that C_4 photosynthesis becomes
479 disadvantageous under a set of environmental conditions (low light, low temperatures and high
480 CO_2) thus providing theoretical support for decades of ecophysiological observations. For the
481 expression of a CCM to be advantageous, any modifications to dark reactions need to be
482 accompanied by substantial modifications to light reactions. Specifically, engineering an
483 appropriate electron transport chain, with the possibility of expressing the NDH complex and
484 adjusting levels of cyclic electron flow will be required. These will also need to be accompanied by
485 anatomical modifications to accommodate the biochemical and light harvesting machinery and by
486 the expression of suitable levels of transporters to allow adequate metabolite traffic.

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498 **Author Contributions**

499 CB conceived of the research, performed measurements, developed and coded the models, ran
500 simulations. CB and GDF wrote the paper.

501 **Availability**

502 The model, coded in Excel, is made freely available in Supporting Information. The model does
503 not include 'live' scripts and is fully operational in the open access suite 'Apache Open Office'.

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Tables.

Table 1. Acronyms, definitions, values, and units used.

Symbol / Acronym	Definition	Values / Units	Source
A	Net assimilation	$\mu\text{mol m}^{-2} \text{s}^{-1}$	output
a_t	Total concentration of adenylates in chloroplast	12.7 mmol m^{-2}	Farquhar and Wong (1984)
BS	Bundle sheath		
C_{BS}	CO_2 concentration in the BS	$\mu\text{mol mol}^{-1}$	output
CCM	Carbon concentrating mechanism		
CEF	Cyclic electron flow		
C_M	CO_2 concentration in the M	$\mu\text{mol mol}^{-1}$	variable, iteratively found
D_S	Leaf to boundary layer water mole fraction gradient in the light ($10 \times \text{VPD}$ in kPa)	10 mmol $\text{H}_2\text{O mol air}^{-1}$	gas exchange
D_{S0}	Leaf to boundary layer water mole fraction gradient in the dark ($10 \times \text{VPD}$ in kPa)	8.6 mmol $\text{H}_2\text{O mol air}^{-1}$	gas exchange
E_t	Total concentration of Rubisco sites	6.7 mmol m^{-2}	adjusted from Farquhar and Wong (1984) by fitting gas exchange data
f_C Rubisco	Parameter defining the fraction of actual Rubisco carboxylation in BS relative to leaf-level	dimensionless	set to equal V_{CBS}/V_C output of the enzyme-limited model
f_{Cyc}	Fraction of J_1 following CEF	dimensionless	fitted to max A
f_{NDH}	Fraction of CEF through the NDH complex	0 (C_3 and C_2); 0.2 and 0± (C_2+C_4); 0.4 and 0± (C_4)	assigned
f_{PPDK}	Parameter defining the fraction of PPDK activity in the BS relative to leaf-level	0	assigned
$f_{\text{PR}}, f_{\text{CS}}$	Parameter defining the fraction of activity in BS relative to leaf-level, of phosphoglycerate reduction rate, and carbohydrate synthesis	0 (variable for the simulations of Figure 6B and 6D)	assigned
$f_{\text{Pseudocyc}}$	Fraction of J_1 following used by alternative sinks of electrons like nitrate reduction and the water-water cycle	0.1	assigned (Yin & Struik, 2012)
f_Q	Fraction of J_1 going through the Q-Cycle	1	Yin and Struik (2012)
f_{RLIGHT}	parameter defining the fraction of respiration in the light in BS relative to leaf-level	0 (C_3); 0.2 (C_2); 0.5 (C_2+C_4 , and C_4)	(von Caemmerer, 1989; von Caemmerer, 2000)
g_{BS}	Bundle sheath conductance to CO_2 diffusion	0.00287† mol $\text{m}^{-2} \text{s}^{-1}$	Yin <i>et al.</i> (2016)
GDC	Glycine decarboxylase		
g_M	Mesophyll conductance to CO_2 diffusion	0.26† mol $\text{m}^{-2} \text{s}^{-1}$	gas exchange
g_{S0}	Stomatal conductance in the dark	0.047 mol $\text{CO}_2 \text{m}^{-2} \text{s}^{-1}$	gas exchange
h	Stoichiometry of ATP synthase: protons required to synthesize ATP	4.67 protons / ATP	Vollmar <i>et al.</i> (2009); Hahn <i>et al.</i> (2018)
$l, l_1, l_2, l_{10}, l_{20}$	Light absorbed by PSI and PSII, by PSI, by PSII, by PSI when $f_{\text{Cyc}}=0$, by PSII when $f_{\text{Cyc}}=0$, respectively	μmol $\text{m}^{-2} \text{s}^{-1}$	output
J_1, J_2	Electron flow through PSI, and PSII, respectively	μmol $\text{m}^{-2} \text{s}^{-1}$	output
J_{ATP}	Total leaf-level ATP production rate	μmol $\text{m}^{-2} \text{s}^{-1}$	output
J_{NADPH}	Total leaf-level NADPH production rate	μmol $\text{m}^{-2} \text{s}^{-1}$	output
J_{SAT}	PPFD saturated electron transport rate	310† μmol $\text{m}^{-2} \text{s}^{-1}$	chlorophyll fluorescence
K_C	Rubisco Michaelis-Menten constant for CO_2 in the liquid phase	8† μM	(Wei <i>et al.</i> , 1994; von Caemmerer, 2000; Galmés <i>et al.</i> , 2016; Hermida-Carrera <i>et al.</i> , 2016)
K_h	Effective hydraulic conductance from the soil to the epidermis	12 mmol $\text{H}_2\text{O m}^{-2} \text{s}^{-1} \text{MPa}^{-1}$	Bellasio <i>et al.</i> (2017)
$K_{h\text{CO}_2}$	volatility of CO_2	30.3† μbar μM ⁻¹	Sander (2015)
$K_{h\text{O}_2}$	volatility of O_2	833.3† μbar μM ⁻¹	Warneck and Williams (2012)
K_O	Rubisco Michaelis-Menten constant for O_2 in the liquid phase	335† μM	von Caemmerer (2000)
M	Mesophyll		
NDH	NAD(P)H Dehydrogenase-like (complex)		
O_M, O_{BS}	O_2 concentration in M cells or BS cells	O_M is 210000 or 21000 μmol mol^{-1} O_{BS} is calculated	O_M is assigned O_{BS} is output
p	Concentration of photophosphorylation sites	7.5 mmol m^{-2}	adjusted from Farquhar and Wong (1984) by fitting gas exchange data
PEPC	Phosphoenolpyruvate carboxylase		
PEPCK	Phosphoenolpyruvate carboxykinase		
PPDK	Pyruvate phosphate dikinase		

<i>PPFD</i>	Photosynthetic photon flux density	$\mu\text{mol m}^{-2} \text{s}^{-1}$	assigned
R_{LIGHT} , $R_{\text{LIGHT BS}}$, $R_{\text{LIGHT M}}$	Respiration in the light, leaf-level, in the BS or in the M, respectively input parameter defining the activity of PEPCK relative to V_P	R_{LIGHT} is $0.8\ddagger \mu\text{mol m}^{-2} \text{s}^{-1}$ 0	gas exchange, 0.5 is the assumed fraction of BS respiration relative to total assigned
f_{PEPCK}	Rubisco		
Rubisco	Ribulose biphosphate carboxylase oxygenase		
RuBP	Ribulose-1,5-bisphosphate		
s	lumped energy conversion coefficient (Yin <i>et al.</i> , 2009)	$0.38 \text{ e}^-/\text{quanta}$	gas exchange
$S_{\text{C/O}}$	Rubisco specificity	$2800\ddagger \mu\text{bar}/\mu\text{bar}$ (gas) or $102\ddagger \text{mol mol}^{-1}$ (liq)	gas exchange
$V_{\text{C MAX}}$, V_{C} , $V_{\text{C BS}}$, $V_{\text{C M}}$	Rubisco carboxylation rate, CO_2 saturated, leaf-level, in the BS, in the M, respectively	$V_{\text{C MAX}}$ is $93\ddagger \mu\text{mol m}^{-2} \text{s}^{-1}$	$V_{\text{C MAX}}$ by gas exchange, or output
V_{O} , $V_{\text{O BS}}$, $V_{\text{O M}}$	Rubisco oxygenation rate, total, in the BS or in the M, respectively	$\mu\text{mol m}^{-2} \text{s}^{-1}$	output
$V_{\text{P(J)}}$	Leaf-level actual phosphoenolpyruvate carboxylation rate, inputs to the light-limited model	0 (C_3 and C_2); $0.075J_{\text{ATP}}(\text{C}_2+\text{C}_4)$; $0.2J_{\text{ATP}}(\text{C}_4) \mu\text{mol m}^{-2} \text{s}^{-1}$ (variable for the simulations in Figure 6)	assigned (von Caemmerer, 2000)
$V_{\text{P MAX}}$	CO_2 saturated phosphoenolpyruvate carboxylation rate	0 (C_3 and C_2); $0.1J_{\text{SAT}}(\text{C}_2+\text{C}_4)$; $0.3J_{\text{SAT}}(\text{C}_4) \mu\text{mol m}^{-2} \text{s}^{-1}$	assigned
V_r	Potential pool size of RuBP	150 mmol m^{-2}	Farquhar and Wong (1984)
$Y(I)_{\text{LL}}$	Yield of PSI extrapolated under zero <i>PPFD</i>	1	Yin and Struik (2012)
$Y(II)_{\text{LL}}$	Yield of PSII extrapolated under zero <i>PPFD</i>	0.75	chlorophyll fluorescence
$Y(II)_{\text{MOD}}$	Yield of PSII modelled empirically by a non-rectangular hyperbola	dimensionless	output
α	parameter scaling BS O_2 evolution to net assimilation attributed to BS activity	1	von Caemmerer (1989)
Y^*	Half the reciprocal Rubisco specificity $\gamma^* = \frac{1}{2S_{\text{C/O}}}$	dimensionless	output
θ	Curvature of the non-rectangular hyperbola used to model the light-dependence of $Y(II)$	0.5	gas exchange
θ_A	Curvature of the non-rectangular hyperbola used to smooth the combination of light-limited and enzyme-limited models	0.95	Buckley <i>et al.</i> (2016)
ξ_{GDC}	Specifies the fraction of V_{OM} which is decarboxylated in the BS.	0 (C_3) and 1 (C_2 , C_2+C_4 , and C_4)	Sage and Khoshravesh (2016)
χ_I	Fraction of I shifting from PSII to PSI upon engagement of CEF	dimensionless	output
χ_{Rubisco}	parameter defining the fraction of $V_{\text{C MAX}}$ in BS relative to leaf-level	dimensionless	fitted to maximise A
$\chi_s\beta$	Parameter lumping turgor to conductance scaling factor and the hydromechanical / biochemical response parameter	$0.055 \text{ mol air mmol}^{-1} \text{ ATP s}^{-1} \text{ MPa}^{-1}$	fitted to gas exchange data
ψ_{Soil} , ψ_{Soil0}	Soil water potential in the light or in the dark respectively	0 MPa	assigned
π_e	Epidermal osmotic pressure	1.2 MPa	Bellasio <i>et al.</i> (2017)

† The value shown is at 25 °C but the quantity was made temperature-dependent; ‡ Alternative scenarios in Figure 4

Figures.

Figure 1. Modelling framework. Blue boxes show inputs while orange boxes show outputs; grey boxes represent submodels. Inputs with a thick blue outline are made temperature-dependent. Submodels contoured in red are originally developed for this work. Photosynthetic photon flux density ($PPFD$) is an input to the electron transport submodel to calculate the total ATP production rate (J_{ATP}) and the total NADPH production rate (J_{NADPH}). These and dummy values for CO_2 concentration at the M carboxylating sites (C_M) are fed into the light- and enzyme-limited submodels (Dashed boxes). The outputs from the photosynthesis submodels are used to calculate chloroplastic ATP concentration (τ) and a smoothed combination of the submodels is fed into a stoichiometric submodel to calculate fluxes and reaction rates. τ is used in the stomatal submodel along with inputs for soil water potential (Ψ_{soil}) and evaporative demand (D_s). The output stomatal conductance (g_s) is used to calculate CO_2 concentration in the sub-stomatal cavity (C_i) from external CO_2 concentration (C_a) and in turn used to calculate C_M , which is iterated. See Table 1 for more abbreviations.

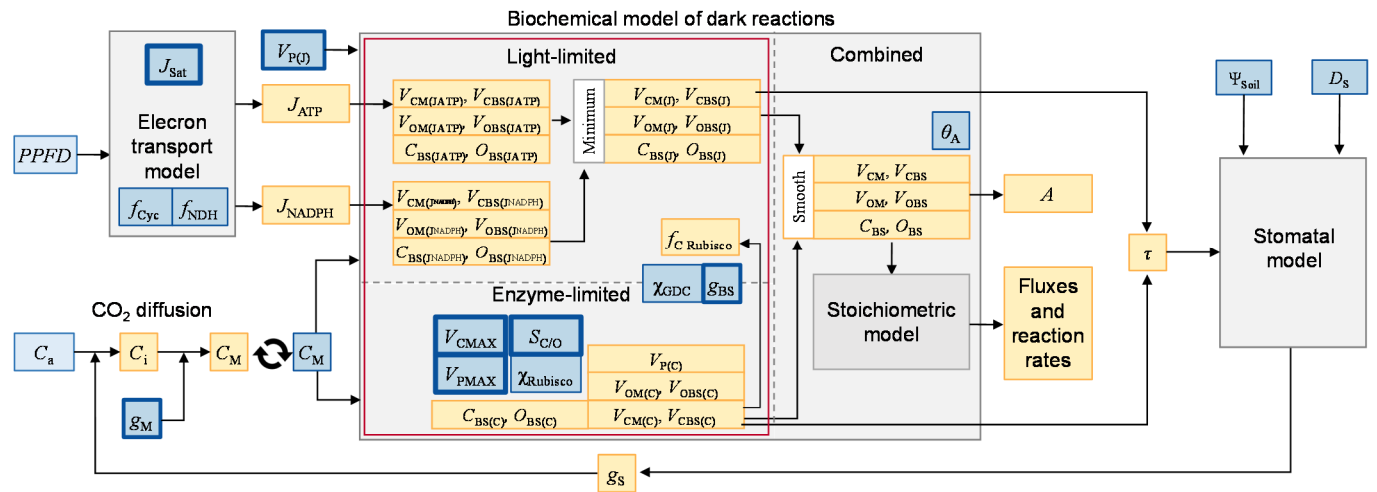


Figure 2. Assimilation and stomatal conductance measured on rice and corresponding simulations for a C_3 photosynthetic type. **Panel A:** light–response curves. Symbols show the response of assimilation (A) to decreasing light intensity ($PPFD$) measured under ambient O_2 (closed circles) or 2% O_2 (open circles). Lines show modelled assimilation under ambient O_2 (solid line) or 2% O_2 (dashed line). **Panel B:** A/C_i curves. Symbols show the measured A at varying levels of CO_2 concentration in the substomatal cavity, C_i , under ambient and low O_2 . Lines show the corresponding simulations. **Panel C:** measured and simulated response of stomatal conductance (g_s) to $PPFD$ under ambient and low O_2 . **Panel D:** measured and simulated response of stomatal conductance (g_s) to external CO_2 concentration, C_a , under ambient and low O_2 . Symbols show mean \pm SE, $n=4$. For simulated A/C_i curves, C_a was set at 16 levels [between 20 and 1000 $\mu\text{mol mol}^{-1}$] while $PPFD$ was set at 1200 $\mu\text{mol m}^{-2} \text{s}^{-1}$, the same used for gas exchange measurements. For simulated $A/PPFD$ curves, $PPFD$ was set at 18 levels [between 1 and 1500 $\mu\text{mol m}^{-2} \text{s}^{-1}$] and C_a was set at 400 $\mu\text{mol mol}^{-1}$. Temperature was 25° C while χ_{Rubisco} and f_{Cyc} were fitted for each combination of inputs.

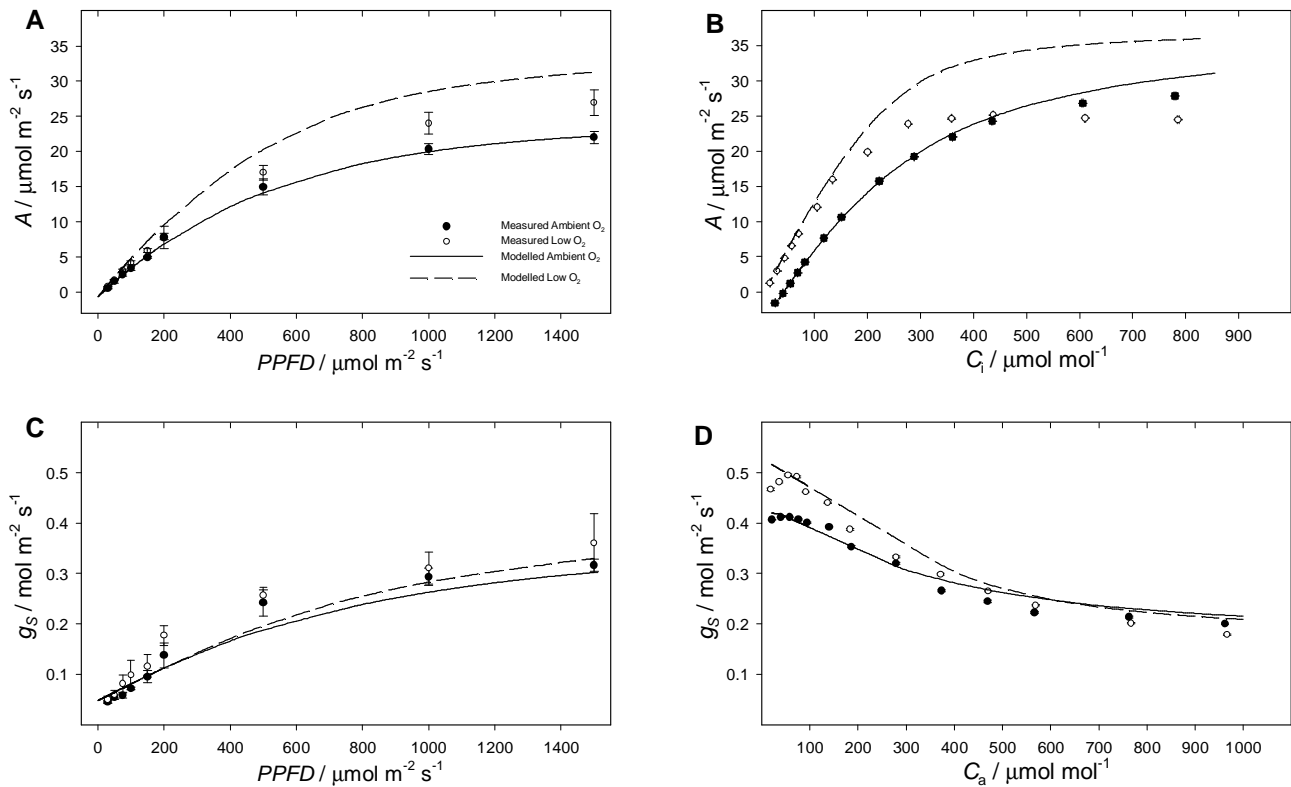


Figure 3. Simulated A -response curves. Compared model output for the four photosynthetic types in response to changes in $PPFD$ (Left) or C_a (Right) varied in the same steps of curves above. Four different photosynthetic types were simulated in a best case scenario for bioengineering whereby the NDH complex is expressed ($f_{NDH} > 0$), f_{Cyc} and $\chi_{Rubisco}$ are optimal (fitted to max A): C_3 (black solid line), representing the measured plants; C_2 (orange dashed line); C_2+C_4 (red solid line); and C_4 (blue dash-dot line). Panels **A** and **B**: net assimilation. Panels **C** and **D**: stomatal conductance. Panels **E** and **F**: CO_2 concentration in the BS. Panels **G** and **H**: Rubisco rate of oxygenation to carboxylation V_o/V_c . Panels **I** and **J**: fraction of Rubisco carboxylating activity in the BS, relative to total.

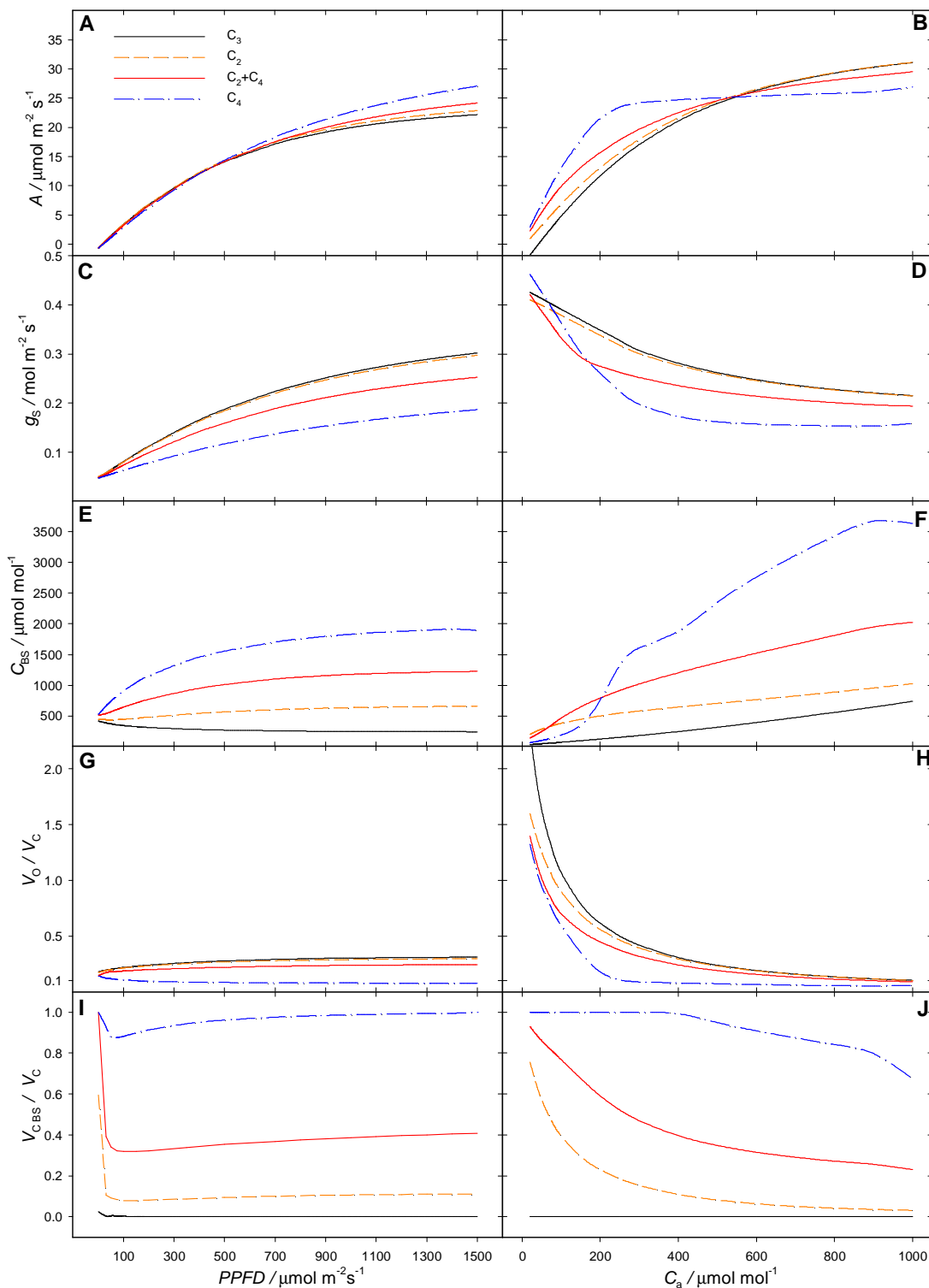


Figure 4. Assimilation in the best case scenario. Gains were calculated for 100 combinations of temperature (varied in 10 steps from 16 °C to 43 °C) and *PPFD* (varied in 10 steps from 1 to 1500 $\mu\text{mol m}^{-2} \text{s}^{-1}$), under a C_a of 400 $\mu\text{mol mol}^{-1}$ (top row), or in 100 combinations of C_a (varied in 10 steps from 150 to 690 $\mu\text{mol mol}^{-1}$) and temperature (as above), under a *PPFD* of 700 $\mu\text{mol m}^{-2} \text{s}^{-1}$ (bottom row) in a best case scenario whereby electron transport processes fully accommodate for the presence of different types of CCM ($f_{\text{NDH}} > 0$, f_{Cyc} is fitted) and Rubisco is optimally allocated (χ_{Rubisco} is fitted). The gain was expressed as relative to C_3 assimilation (Panels **A** and **E**), for C_2 (Panel **B** and **F**) C_2+C_4 (panel **C** and **G**) and C_4 (panel **D** and **H**).

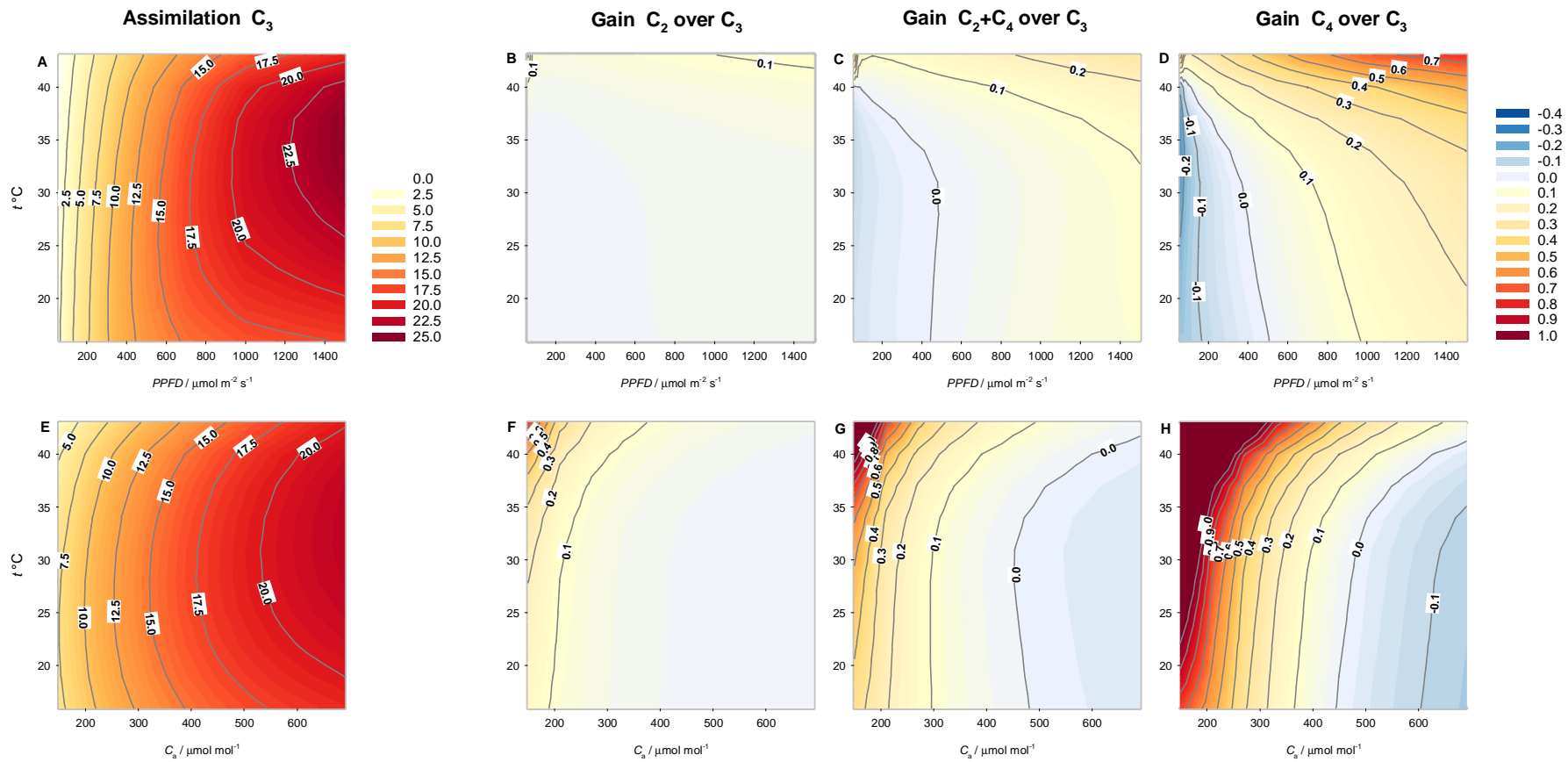


Figure 5. Assimilation in alternative scenarios. Gains were calculated in the temperature \times *PPFD* space, under a C_a of $400 \mu\text{mol mol}^{-1}$, expressed as relative to C_3 assimilation (Figure 4). Panels **A**, **B**, and **C** show a less optimistic scenario whereby the activity of the NDH complex remain at C_3 levels, modelled by setting f_{NDH} at zero for all photosynthetic types. Panels **D**, **E** and **F** show a pessimistic scenario whereby in addition to $f_{\text{NDH}}=0$, the fraction of cyclic electron flow (f_{Cyc}) was set at C_3 levels for all photosynthetic types.

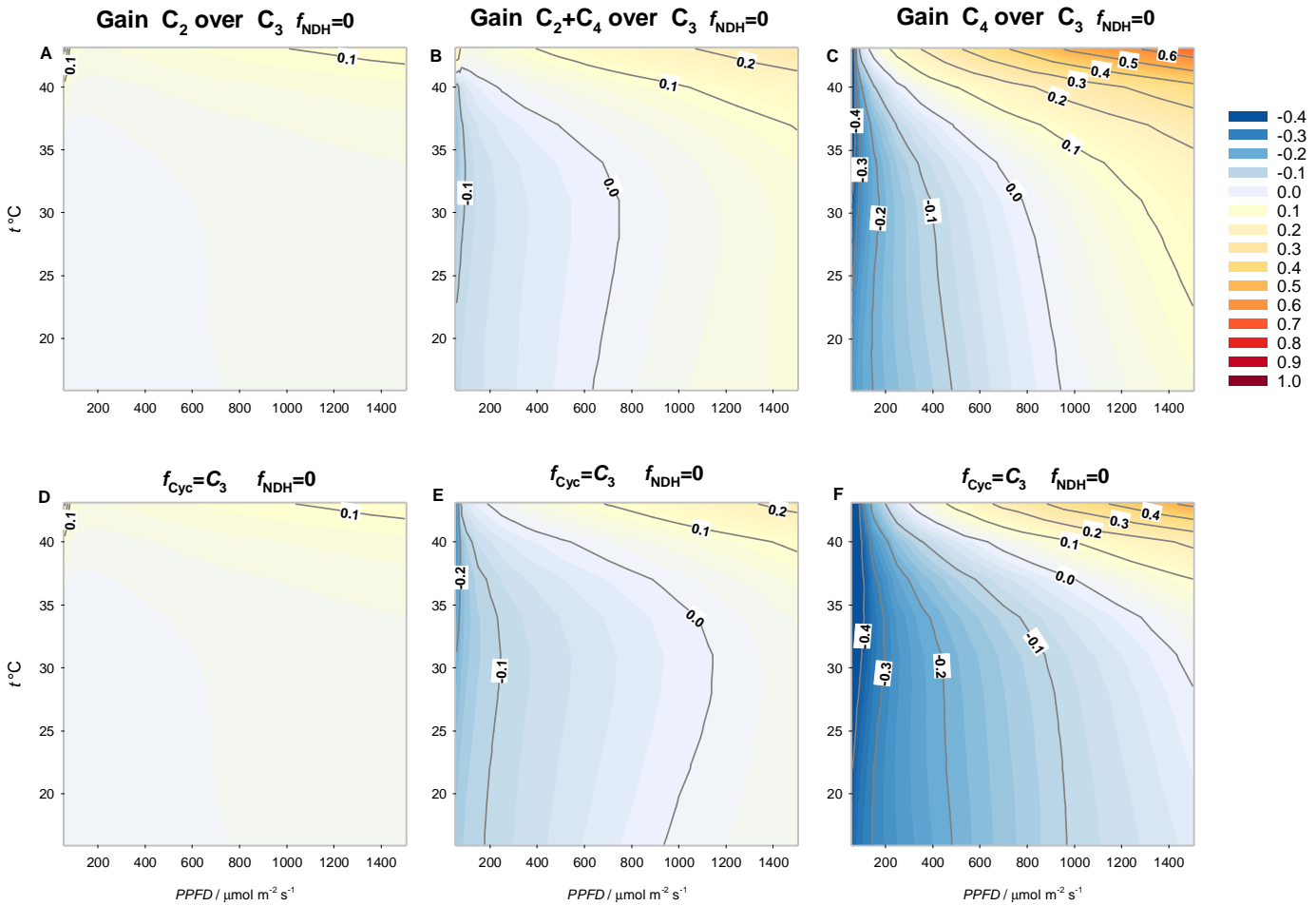


Figure 6. Modelled fluxes between the M and the BS at increasing levels of C₄ engagement. In this simulation the C₄ CCM was increasingly upregulated by manipulating PEPC activity ($V_{P(J)}$, $\mu\text{mol m}^{-2} \text{s}^{-1}$) to increase from 0 to 0.2 J_{ATP} to represent the C₂ to C₄ continuum (from left to right of each panel). Panel **A** simulates a scenario of minimum ATP demand in BS obtained by setting r_{PEPCK} , f_{PR} , f_{CS} and f_{PPDK} at zero; other inputs represented the operational conditions of $PPFD$ 700 $\mu\text{mol m}^{-2} \text{s}^{-1}$, 25 °C, and $C_a=350 \mu\text{mol mol}^{-1}$. Panel **B** simulates a scenario of minimum sum of squared flow rates between BS and M obtained by fitting f_{PR} , and f_{CS} . In these conditions the ATP demand in BS increased substantially, and is shown as relative to the ATP demand in the M in panels **C** and **D**. The flux is considered positive when in the M to BS direction for MAL, ASP, DHAP, and GLY, and in the opposite direction for the other metabolites (Figure S1). Note the different scaling of y-axes.

