

1 Synthetic biology approaches for improving photosynthesis

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9 Keywords: carbon fixation, Rubisco, Calvin Cycle, photorespiration, carbon concentrating  
10 mechanisms, C4 metabolism

11 **Highlight section**

12 We discuss current efforts to boost plant carbon fixation using a synthetic biology approach,  
13 highlighting the engineering of Rubisco, optimizing the Calvin Cycle, introducing carbon  
14 concentrating mechanisms, and rewiring photorespiration.

## 15 **Abstract**

16 The phenomenal increase in agricultural yields that we have witnessed in the last century has  
17 slowed down as we approach the limits of selective breeding and optimization of cultivation  
18 techniques. To support the yield increase required to feed an ever growing population, we will have  
19 to identify new ways to boost the efficiency by which plants convert light into biomass. This  
20 challenge could be potentially tackled using state-of-the-art synthetic biology techniques to rewrite  
21 plant carbon fixation. In this review, we use recent studies to discuss and demonstrate different  
22 approaches for enhancing carbon fixation, including engineering Rubisco for higher activity,  
23 specificity, and activation; changing the expression level of enzymes within the Calvin Cycle to  
24 avoid kinetic bottlenecks; introducing carbon concentrating mechanisms such as inorganic carbon  
25 transporters, carboxysomes, and C4 metabolism; and rewiring photorespiration towards more  
26 energetically efficient routes or pathways that do not release CO<sub>2</sub>. We conclude by noting the  
27 importance of prioritizing and combining different approaches towards continuous and sustainable  
28 increase of plant productivities.

## 29 Introduction

30 Selective breeding and optimization of cultivation techniques have historically driven increases in  
31 agricultural output. In the last century, these efforts have adopted a more scientific approach with  
32 the development of the Haber-Bosch process (Haber and Le Rossignol, 1909; Sutton *et al.*, 2008),  
33 and, later, the “green revolution” (Khush, 2001). Since 1961, global rice and wheat yields increased  
34 by 150% and 210%, respectively (FAO, 2018). However, we have recently started to witness  
35 stagnation in growth improvement of major crops such as rice in China (Peng *et al.*, 2009) or wheat  
36 in the USA (Ray *et al.*, 2012). This presents a major problem, as further yield increases are sorely  
37 needed to feed human population, especially considering global shift towards meat-dependent  
38 diets, use of arable lands to feed bio-refineries, deleterious effects of climate change, and  
39 continuous erosion of agricultural land (Godfray *et al.*, 2010; Tilman *et al.*, 2011).

40 Agricultural yield can be modeled as a product of three factors (Monteith and Moss, 1977; Fletcher  
41 *et al.*, 2011): (i) efficiency of intercepting light; (ii) efficiency of converting intercepted light into  
42 biomass; and (iii) the harvest index, i.e., the fraction of biomass that is captured in the harvested  
43 part. In the past, improved yields have largely been achieved by increasing the light capture  
44 efficiency and the harvest index; however, these two factors now appear to approach their practical  
45 limits (Long *et al.*, 2006). Therefore, the efficiency by which plants convert light to biomass has  
46 become the prime focus for further improvement (Long *et al.*, 2006). This efficiency is determined by  
47 two main processes, the light-dependent reactions, in which photoenergy is used for the generation  
48 of the cellular redox and energy carriers NADPH and ATP, and the light-independent reactions,  
49 which use these carriers to fix CO<sub>2</sub> and reduce it to organic carbon. The efficiency of both processes  
50 is unlikely to be improved by a classic selective breeding approach – as demonstrated by a recent  
51 study exploring 80 years of soybean breeding (Koester *et al.*, 2016) – but could be potentially  
52 increased by dedicated engineering (Zhu *et al.*, 2010). The focus of this review is the use of  
53 synthetic biology tools for boosting the efficiency and rate of carbon fixation. Rather than discuss  
54 the technical aspects of synthetic biology in plants – for which we refer the readers to other reviews  
55 (DePaoli *et al.*, 2014; Liu and Stewart, 2015; Boehm and Bock, 2018; Piatek *et al.*, 2018; Vazquez-  
56 Vilar *et al.*, 2018) – we emphasize conceptual strategies to boost carbon fixation. In particular, we  
57 discuss efforts aiming to improve carboxylation by Rubisco, optimize expression levels of enzymes  
58 within the Calvin Cycle, introduce carbon concentration mechanisms, and rewire photorespiration.  
59 We claim that multiple complementary strategies are paving the way towards substantial yield  
60 increases that are not feasible using conservative selective breeding techniques.

## 61 **Engineering Rubisco**

62 Rubisco, the key enzyme of the Calvin cycle, is probably the most abundant protein in the biosphere  
63 (Ellis, 1979; Raven, 2013), and is responsible for assimilating the vast majority of inorganic carbon  
64 (Raven, 2009). The enzyme catalyzes the condensation of ribulose 1,5-bisphosphate (RuBP) with  
65 CO<sub>2</sub> to give two molecules of glycerate 3-phosphate (G3P). Despite its key biochemical role,  
66 Rubisco is considerably slower than most enzymes in central metabolism (Bar-Even *et al.*, 2011).  
67 Moreover, Rubisco is not completely specific to CO<sub>2</sub> and also accepts O<sub>2</sub>, leading to the formation of  
68 2-phosphoglycolate (2PG) that needs to be reassimilated. In the C<sub>3</sub> model plant *A. thaliana*, the  
69 carboxylation to oxygenation ratio was measured to be as low as 2.3:1 at high light conditions (Ma  
70 *et al.*, 2014). Suppressing oxygenation reactions by cultivating plants at elevated CO<sub>2</sub>  
71 concentrations has repeatedly shown to increase productivity. For example, a meta-analysis of 70  
72 studies showed that rice yields increased by 23% when CO<sub>2</sub> concentrations were raised to 627 ppm  
73 (Ainsworth, 2008). These results indicate that engineering Rubisco for higher CO<sub>2</sub> specificity could  
74 substantially boost yield.

75 Approaches to improve Rubisco catalysis by random or site-directed mutagenesis have generally  
76 failed to yield substantial kinetic enhancements (Somerville and Ogren, 1982; Spreitzer *et al.*, 2005;  
77 Whitney *et al.*, 2011; Wilson *et al.*, 2016). Comparisons between Rubisco variants from a range of  
78 different organisms have revealed a trade-off between CO<sub>2</sub> specificity and carboxylation velocity  
79 (Tcherkez *et al.*, 2006; Savir *et al.*, 2010; Galmés *et al.*, 2014), although several recent studies  
80 challenge this finding (Young *et al.*, 2016; Cummins *et al.*, 2018). Considering this tradeoff, it  
81 actually seems that most Rubisco variants are well adapted to their intracellular environment. Still,  
82 as ambient CO<sub>2</sub> concentrations are changing at a rate faster than plants can adapt to, it was  
83 suggested that replacing plant Rubisco with another variant could boost carbon fixation by up to  
84 25% (Zhu *et al.*, 2004; Orr *et al.*, 2016). Substituting one Rubisco variant with another is  
85 undoubtedly a challenging task, but was already demonstrated using homodimeric Rubisco from the  
86  $\alpha$ -proteobacterium *R. rubrum* (Whitney and Andrews, 2001) and, more recently, using a fast  
87 hexadecameric Rubisco from *S. elongates* (Lin *et al.*, 2014; Occhialini *et al.*, 2016). Coexpression of  
88 supporting chaperones, including the appropriate accumulation factors, can assist in producing an  
89 active Rubisco recombinantly, and can further facilitate efforts to enhance the kinetics of this key  
90 enzyme via mutagenesis (Aigner *et al.*, 2017).

91 Carbon fixation via Rubisco can be potentially improved by means other than direct engineering of  
92 its catalytic parameters. The addition of a CO<sub>2</sub> molecule to an active site lysine, i.e., carbamylation,  
93 is a prerequisite for Rubisco activity (Lorimer and Miziorko, 1980), but can be hindered by the  
94 premature binding of RuBP or other sugar phosphates (Portis, 2002; Parry *et al.*, 2007). The

95 catalytic chaperone Rubisco activase (Rca) removes the sugar phosphate inhibitors from an inactive  
96 uncarbamylated enzyme or an inhibited carbamylated Rubisco (Portis, 2002). As the thermal  
97 instability of Rca was shown to constrain carbon fixation under moderate heat stress (Salvucci *et al.*,  
98 2004), it has become an attractive target for engineering towards enhanced photosynthesis. For  
99 example, by increasing the thermostability of Rca in *A. thaliana*, improved photosynthesis and  
100 growth rate were demonstrated under a moderate heat stress (Kurek *et al.*, 2007; Kumar *et al.*,  
101 2009). Similarly, overexpression of maize Rca in rice led to higher activation state of Rubisco in low  
102 light and faster response of photosynthesis when light intensities increased (Yamori *et al.*, 2012).

### 103 **Optimizing expression of Calvin Cycle enzymes**

104 G3P produced by Rubisco need to be metabolized by nine enzymes of the Calvin cycle to  
105 regenerate RuBP. This regeneration process, whose rate has to match that of Rubisco, is known to  
106 limit carbon fixation rate under certain conditions. Computational models have suggested that the  
107 natural distribution of enzymes within the Calvin Cycle is not optimal and could limit photosynthesis  
108 (Zhu *et al.*, 2007). Specifically, it was predicted that higher levels of sedoheptulose-1,7-  
109 bisphosphatase and fructose-1,6-bisphosphate aldolase, as well as enzymes linked to sink  
110 capacity, could support higher productivity.

111 Unsurprising, under elevated CO<sub>2</sub> concentrations, the rate of Rubisco becomes less limiting and  
112 carbon fixation is mostly constrained by RuBP regeneration. For example, studies of *N. tabacum* at  
113 930 ppm CO<sub>2</sub> showed that reducing Rubisco levels by 30-50% did not inhibit growth (Masle *et al.*,  
114 1993). Similar results were obtained in rice plants in which Rubisco levels were reduced by 65% at  
115 1000 ppm CO<sub>2</sub>. On the other hand, overexpression of sedoheptulose-1,7-bisphosphatase in *N.*  
116 *tabacum* at 585 ppm CO<sub>2</sub> resulted in higher carbon fixation rate (Rosenthal *et al.*, 2011). Similarly,  
117 at 700 ppm, increased levels of fructose-1,6-bisphosphate aldolase in *N. tabacum* led to increased  
118 biomass (Uematsu *et al.*, 2012).

119 Even at ambient CO<sub>2</sub> concentration, overexpression of limiting enzymes of the Calvin Cycle was  
120 shown to boost carbon fixation. In *N. tabacum*, overexpression of sedoheptulose 1,7-  
121 bisphosphatase (Lefebvre *et al.*, 2005) and fructose 1,6-bisphosphatase (Miyagawa *et al.*, 2001)  
122 increased photosynthetic rates and biomass. Similarly, the co-overexpression of sedoheptulose-1,7-  
123 bisphosphatase and fructose-1,6-phosphate aldolase enhanced photosynthesis and yield (Simkin *et*  
124 *al.*, 2015).

## 125 **Establishing carbon concentrating mechanisms**

126 To mitigate the problem of oxygenation, and further enable the use of faster (and less specific)  
127 Rubisco, multiple organisms have developed carbon concentrating mechanisms (CCMs) to  
128 concentrate CO<sub>2</sub> at the site of Rubisco. As C<sub>3</sub> plants lack CCMs, it was proposed to introduce them  
129 to increase photosynthetic efficiency. Two main approaches are actively pursued: (A) introduction of  
130 biophysical CCMs from cyanobacteria and green algae (Long *et al.*, 2016; Rae *et al.*, 2017); and (B)  
131 introduction of C<sub>4</sub> anatomy and metabolism (Hibberd *et al.*, 2008; Schuler *et al.*, 2016).

132 Biophysical CCM are found in cyanobacteria (Kupriyanova *et al.*, 2013) and in green algae like *C.*  
133 *reinhardtii* (Mackinder, 2018). In such CCM, bicarbonate is actively transported into the cytosol in  
134 which carbonic anhydrase is lacking. From there, bicarbonate is further transported into specialized  
135 compartments packed with Rubisco – carboxysomes in cyanobacteria and pyrenoids in green algae  
136 – where it is dehydrated to CO<sub>2</sub> by carbonic anhydrase. It is thought that both carboxysomes and  
137 pyrenoids present a diffusion barrier for CO<sub>2</sub> and O<sub>2</sub>, keeping the former molecule in and the latter  
138 molecule out, and thus enhancing carboxylation and suppressing oxygenation (Mangan *et al.*,  
139 2016).

140 Establishing biophysical CCM in plants is a challenging task that first requires the expression and  
141 correct localization of inorganic carbon transporters. It was suggested that the transporters  
142 themselves could increase carbon fixation rate albeit to a limited extent (McGrath 2014, Yin 2017).  
143 Indeed, overexpression of the putative-inorganic carbon transporter from cyanobacteria, *ictB*, in *A.*  
144 *thaliana*, tobacco, rice, and soybean was reported to increase photosynthetic rate and biomass  
145 (Lieman-Hurwitz *et al.*, 2003, 2005; Yang *et al.*, 2008; Simkin *et al.*, 2015; Hay *et al.*, 2017). In  
146 contrast, expression of other transporters from cyanobacteria or *C. reinhardtii* did not increase yield  
147 or improve growth, despite correct localization within the plant cells (Rolland *et al.*, 2016; Uehara *et*  
148 *al.*, 2016; Atkinson *et al.*, 2016). Optimizing transporter activity is therefore still an open challenge  
149 that needs to be resolved before commencing with the next step: assembly of Rubisco containing  
150 compartments. The establishment of these sophisticated structures would enable further increase in  
151 CO<sub>2</sub> concentration at the site of Rubisco and could therefore substantially enhance carbon fixation.  
152 Recently, simplified carboxysome structures were introduced into the chloroplasts of *N. tabacum*  
153 (Long *et al.*, 2018). Yet, these are expected to enhance photosynthesis only after combination with  
154 functional inorganic carbon transporters (McGrath and Long, 2014).

## 155 **Engineering C4 metabolism**

156 As an alternative to biophysical CCM, ongoing research is dedicated to introducing C4 metabolism  
157 into C3 plants (Schuler *et al.*, 2016). C4 metabolism utilizes the most efficient carbon fixation  
158 enzyme – PEP carboxylase – to temporarily capture inorganic carbon, which is then transported to  
159 the vicinity of Rubisco (Jenkins *et al.*, 1989). Specifically, PEP carboxylase in the mesophyll cells  
160 ‘borrows’ PEP and converts it to oxaloacetate, which is further metabolized to malate or aspartate.  
161 These C4 acids are transported to the bundle sheath cells and decarboxylated to release CO<sub>2</sub> next  
162 to RubiCO, which is mainly localized in these cells. Pyruvate, the product of this decarboxylation, is  
163 then transported back to the mesophyll cells to regenerate PEP. Hence, the entire C4 cycle, which  
164 depends on a special anatomy termed “Kranz anatomy” (mesophyll cells surrounding bundle sheath  
165 cells), can be regarded as a sophisticated CO<sub>2</sub> pump that results in ~10 times higher concentration  
166 of inorganic carbon in the vicinity of RubiCO (Jenkins *et al.*, 1989).

167 Engineering C4 photosynthesis in C3 plants has been outlined as a stepwise process (Schuler *et*  
168 *al.*, 2016) that includes alteration of plant tissue anatomy, establishment of bundle sheath  
169 morphology, as well as ensuring a cell-type specific enzyme expression. Although challenging,  
170 engineering a C3 plant to have C4 metabolism seems to be a feasible goal as it is known to have  
171 emerged independently at least 66 times in different phylogenetic backgrounds (Sage *et al.*, 2012).  
172 Importantly, C3 plants already harbor the main enzymes of C4 metabolism, e.g., PEP carboxylase  
173 (Aubry *et al.*, 2011), and are known to shuttle carbon from the vasculature into the surrounding cells  
174 in a way similar to that of C4 plants (Hibberd and Quick, 2002; Brown *et al.*, 2010). This provides a  
175 solid basis to replicate the emergence of C4 metabolism by direct engineering.

176 Nevertheless, despite international efforts, a synthetic C4 plant has yet to be reported. Following  
177 Richard Feynman’s famous quote “What I cannot create, I do not understand”, it seems that  
178 incomplete understanding of C4 metabolism hampers its engineering. Specifically, the metabolic  
179 shuttling of intermediates between mesophyll and bundle sheath cells and the factors necessary to  
180 create Kranz anatomy are still not fully clear and need to be elucidated (Schuler *et al.*, 2016).

181 It might not be necessary to establish a complete C4 metabolism in order to improve carbon fixation.  
182 It was recently suggested that engineering a C3-C4 intermediate metabolism could enhance  
183 productivity (Schlüter and Weber, 2016). For example, in C3-C4 intermediate type I plants,  
184 photorespiratory glycine is transported from the mesophyll cells to the bundle sheath cells for  
185 decarboxylation. In the bundle sheath cells, the mitochondria are closely associated with the  
186 chloroplast, thereby enhancing re-assimilation of released CO<sub>2</sub> by nearby RubiCO (Monson and  
187 Edwards, 1984; Rawsthorne *et al.*, 1988). Establishing this intermediary metabolism within C3

188 plants, besides being useful on its own to boost carbon fixation, would provide a milestone towards  
189 further engineering of complete C4 metabolism.

190 An interesting alternative engineering target is crassulacean acid metabolism (CAM). While C4  
191 metabolism increases CO<sub>2</sub> concentration at the vicinity of Rubisco via spatial decoupling, CAM  
192 accomplishes the same goal via temporal decoupling. Specifically, inorganic carbon is temporarily  
193 fixed by the highly efficient PEP carboxylase during the night, when the stomata are open and CO<sub>2</sub>  
194 can freely enter the cell. Malate, the indirect product of the carboxylation, is stored within the  
195 vacuole. During the day, when the stomata are closed, malate is decarboxylated, releasing CO<sub>2</sub> and  
196 maintaining its high concentration for subsequent fixation by Rubisco and the Calvin Cycle. Besides  
197 increasing CO<sub>2</sub> concentration in the vicinity of Rubisco, CAM reduces water evaporation and  
198 increase water-use efficiency by 20-80% (Borland *et al.*, 2009), making CAM plants highly suitable  
199 for arid climates. Similarly to C4 metabolism, CAM has arisen multiple times in a taxonomically  
200 diverse range of plants, indicating that its necessary components exist in C3 plants which could  
201 potentially be engineered towards this unique carbon metabolism (DePaoli *et al.*, 2014).  
202 Furthermore, the existence of C3-CAM intermediate species and plants that switch between both  
203 metabolic modes further supports the potential of engineering C3 metabolism towards CAM  
204 (Borland *et al.*, 2011). Such engineering would require precise control of the activity key enzymes  
205 (e.g., PEP carboxylase, malic enzyme, and Rubisco), stomatal conductance, and intracellular  
206 transport (e.g., to and from the vacuole) (Borland *et al.*, 2014; DePaoli *et al.*, 2014; Yang *et al.*,  
207 2015).

## 208 **Rewiring photorespiration**

209 2PG, the product of Rubisco's oxygenation activity, is recycled to the Calvin Cycle in a process  
210 termed photorespiration. This rather long pathway requires the shuttling of metabolites across  
211 multiple organelles and is considered inefficient as it dissipates energy by releasing ammonia and  
212 using oxygen as an electron acceptor. Moreover, photorespiration releases one CO<sub>2</sub> molecule in the  
213 recycling of two 2PG molecules and hence directly counteracts carbon fixation by the Calvin Cycle.  
214 The inefficiencies associated with the recycling of 2PG cannot be prevented by simply blocking  
215 photorespiration, as this pathway plays an essential role in plant metabolism (Somerville and Ogren,  
216 1979) and reduction of its flux was shown to negatively affect photosynthesis (Servaites and Ogren,  
217 1977; Wingler *et al.*, 1997; Heineke *et al.*, 2001). One explanation for this lies in the inhibitory  
218 effects exerted by several photorespiratory intermediates. For example, 2PG was shown to inhibit  
219 triosephosphate isomerase and sedoheptulose 1,7-bisphosphate phosphatase (Anderson, 1971;  
220 Flügel *et al.*, 2017), glyoxylate impairs Rubisco activation (Chastain and Ogren, 1989; Campbell and  
221 Ogren, 1990; Hausler *et al.*, 1996; Savir *et al.*, 2010), and glycine interferes with Mg<sup>2+</sup> availability

222 (Eisenhut *et al.*, 2007). Based on these observations, it was suggested that increased  
223 photorespiratory flux could prevent the accumulation of inhibitory intermediates and enhance  
224 photosynthesis; indeed, this was demonstrated upon overexpression of components of the glycine  
225 cleavage system in *A. thaliana* (Timm *et al.*, 2012, 2015) and in *N. tabacum* (Lopez-Calcano *et al.*,  
226 2018).

227 While photorespiration cannot be avoided, it might be possible to replace the natural pathway with  
228 more efficient alternatives. The first bypass suggested in this regard was inspired by cyanobacterial  
229 photorespiration (Eisenhut, 2006), where glyoxylate is condensed and reduced to directly generate  
230 the key photorespiratory intermediate glycerate. This pathway was implemented in *A. thaliana*  
231 (Kebeish *et al.*, 2007) and later in *C. sativa* (Dalal *et al.*, 2015) using glycolate dehydrogenase,  
232 glyoxylate carboxyligase, and tartronic semialdehyde reductase from *Escherichia coli*. In both  
233 cases, this metabolic bypass, dissipating less energy and shifting CO<sub>2</sub> release from the  
234 mitochondria to the chloroplast, was shown to increase photosynthesis and biomass.

235 However, it was shown that expression of only the first enzyme of the pathway, glycolate  
236 dehydrogenase, suffices to enhance photosynthesis. Supporting this, chloroplastic expression of  
237 glycolate dehydrogenase in *S. tuberosum* induced a 2.3-fold increase in tuber yield (Nölke *et al.*,  
238 2014). This suggests that the benefits of glycerate-pathway might not stem from more efficient  
239 recycling of 2PG but rather from oxidation of glycolate to glyoxylate. Indeed, incubation with  
240 glyoxylate was shown to increase carbon fixation – potentially due to suppression of Rubisco  
241 oxygenation – in both tobacco leaf disks (Oliver and Zelitch, 1977) and soybean mesophyll cells  
242 (Oliver, 1980).

243 Another photorespiratory bypass involves the complete oxidation of 2PG to CO<sub>2</sub> via a catabolic  
244 pathway that consist of glycolate dehydrogenase, malate synthase, malic enzyme, and pyruvate  
245 dehydrogenase (Maier *et al.*, 2012). While the authors reported increased biomass and  
246 photosynthesis, it is still unclear which mechanism supports the beneficial effect of the pathway, as  
247 a theoretical model predicts a negative effect when 2PG is completely oxidized (Xin *et al.*, 2015).

## 248 **Carbon-conserving photorespiration**

249 As the main problem associated with photorespiration is (arguably) the release of CO<sub>2</sub>, bypasses  
250 that do not lead to the loss of carbon could dramatically boost carbon fixation. Several synthetic  
251 carbon-conserving bypasses have been suggested. In the *de novo* 2PG salvage pathway (Ort *et al.*,  
252 2015), 2PG was suggested to be reduced to 2-phosphoglycolaldehyde, which is subsequently  
253 condensed with dihydroxyacetone phosphate to give xylulose biphosphate. This intermediate is

254 then dephosphorylated to xylulose 5-phosphate, a Calvin cycle metabolite. The main challenges of  
255 this proposed bypass is the reversibility of most of its reactions (resulting in low driving force), the  
256 low concentration of 2PG, and the inhibitory effect of xylulose biphosphate (Yokota, 1991; Zhu and  
257 Jensen, 1991; Parry *et al.*, 2007).

258 Recently, a systematic analysis identified multiple synthetic routes that can bypass photorespiration  
259 without the release of CO<sub>2</sub>. Several of these pathways involve the reduction of glycolate (the  
260 concentration of which is considerably higher than that of 2PG) to glycolaldehyde, which then  
261 undergoes an aldol condensation with a phosphosugar from the Calvin Cycle to generate a longer  
262 chain phosphosugar that is reintegrated into Calvin Cycle (Bar-Even, 2018; Trudeau *et al.*, 2018). A  
263 computational model indicated that these pathways can boost photosynthesis under all  
264 physiologically relevant irradiation and intracellular CO<sub>2</sub> levels.

265 The operation of these carbon conserving bypass routes depends on the conversion of glycolate to  
266 glycolaldehyde, but this activity is not supported by any known enzyme. To establish this activity two  
267 enzymes were engineered (Trudeau *et al.*, 2018). First, acetyl-CoA synthetase from *E. coli* was  
268 engineered to accept glycolate, thus generating glycolyl-CoA. Next, propionyl-CoA reductase from  
269 *Rhodospseudomonas palustris* was engineered to accept glycolyl-CoA, reducing it to glycolaldehyde.  
270 The cofactor specificity of this latter enzyme was switched, such that it could use NADPH – the  
271 photosynthetic electron carrier – as an electron donor. The two engineered enzymes were  
272 combined, in a test-tube, with fructose 6-phosphate aldolase (condensing glycolaldehyde with  
273 glyceraldehyde 3-phosphate to generate arabinose 5-phosphate), arabinose 5-phosphate  
274 isomerase, and phosphoribulokinase. Upon addition of glycolate and glyceraldehyde 3-phosphate,  
275 NADPH and ATP were consumed and RuBP was found to accumulate (Trudeau *et al.*, 2018),  
276 demonstrating the *in vitro* activity of an alternative photorespiration route that does not release CO<sub>2</sub>.

277 It was further proposed to go beyond carbon conservation, and engineer a photorespiration bypass  
278 that fixes CO<sub>2</sub> and thus directly support the activity of the Calvin Cycle. One such a carbon-positive  
279 bypass was inspired by the 3-hydroxypropionate bicycle (Shih *et al.*, 2014). Here, glycolate is  
280 oxidized to glyoxylate, which is then metabolized and further carboxylated to pyruvate. Towards the  
281 implementation of this bypass, six non-native genes from *C. aurantiacus* were expressed in  
282 cyanobacteria, but no distinct growth phenotype was evident.

283 In another study, glycolate was not recycled to the Calvin cycle but instead metabolized to acetyl-  
284 CoA via the synthetic malyl-CoA-glycerate pathway (Yu *et al.*, 2018). This pathway can further be  
285 used to generate acetyl-CoA from photosynthetic C3 sugars via an additional CO<sub>2</sub>-fixing step,

286 thereby bypassing CO<sub>2</sub> release by pyruvate dehydrogenase. In cyanobacteria, the pathway  
287 facilitated a two-fold increase in bicarbonate assimilation.

## 288 **Conclusions**

289 The increasing number of studies demonstrating improved photosynthesis and growth by  
290 engineering different components of the light-dependent and independent reactions indicates that  
291 we are on the right path. Yet, many challenges are ahead of us. Beside the technical difficulties,  
292 which we did not discuss here and refer the reader to other reviews (DePaoli *et al.*, 2014; Liu and  
293 Stewart, 2015; Boehm and Bock, 2018; Piatek *et al.*, 2018; Vazquez-Vilar *et al.*, 2018), there is one  
294 key barrier that worth elaborating on, which is system complexity. Complex systems are notoriously  
295 difficult to engineer as the effect of even small changes can have substantial results that cannot be  
296 easily predicted. While mathematical models can help deal with such complexity, the lack of  
297 knowledge regarding many of involved components commonly hinders accurate prediction. Plant  
298 carbon metabolism provides an excellent example of a complex system, the response of which to  
299 changes is hard to foretell. Previous attempts to engineer carbon fixation demonstrate this vividly.  
300 Perhaps the best example is the engineering of photorespiration bypass routes as described above.  
301 While few bypasses were already shown to enhance photosynthesis, the cause of this effect is  
302 probably different than that originally suggested, as chloroplastic oxidation of glycolate suffices to  
303 support most of the beneficial effects. Unraveling this mystery would require deep understanding of  
304 the intricate interplay between all system components, a task which we have yet to fully achieve.

305 Moreover, while some engineering efforts show only minor benefits in isolation, the key for future  
306 improvements lies in the correct combination of multiple strategies. Indeed, first examples for  
307 beneficial cumulative effects have been reported (Simkin *et al.*, 2015). It is further clear that not all  
308 strategies can be implemented with similar ease. Overexpressing a Calvin Cycle enzyme, for  
309 example, is considerably easier than rerouting photorespiration via a synthetic pathway that does  
310 not release CO<sub>2</sub>. It is therefore important to carefully choose targets for the near- and medium-future  
311 and progress in a way that ensures intermediate gains. For example, establishing a C3-C4  
312 intermediate metabolism does not only provide a solid basis for further engineering of a complete  
313 C4 metabolism, but is expected to boost carbon fixation by itself. Once we gain the required  
314 proficiency in rewiring plant central metabolism, we can aim at even bigger targets, for example,  
315 replacing Rubisco with a set of enzymes, each responsible for a different catalytic step (Bar-Even,  
316 2018), or replacing the Calvin Cycle with a synthetic carbon fixation pathway (Schwander *et al.*,  
317 2016).

318 **Acknowledgements**

319 The authors thank Charlie Cotton for critical reading of the manuscript. This work was funded by the  
320 Max Planck Society and by the European Union's Horizon 2020 FET Programme under the grant  
321 agreement No 686330 (FutureAgriculture).

## **Key developments box**

### **Assembly of Rubisco-containing carboxysomes in tobacco chloroplasts**

Assembly of a simplified  $\alpha$ -carboxysome in tobacco chloroplasts by replacing native Rubisco with large and small subunits of Rubisco from cyanobacteria and two key structural subunits. The introduction of carboxysomes to plant chloroplasts is a key step towards establishing a full biophysical carbon concentrating mechanism in higher plants (Long *et al.*, 2018).

### **Design and *in vitro* realization of carbon-conserving photorespiration**

A systematic search and analysis of synthetic photorespiration bypass routes that do not release CO<sub>2</sub> reveals that these can enhance carbon fixation rate under all relevant physiological conditions. Two enzymes were engineered to jointly enable the reduction of glycolate to glycolaldehyde. The combination of these evolved enzymes with existing ones supported the *in vitro* recycling of glycolate to RuBP without the loss of CO<sub>2</sub>, indicating the feasibility of carbon conserving photorespiration (Trudeau *et al.*, 2018).

### **The synthetic methyl-CoA-glycerate pathway supports photosynthesis**

An *in vivo* demonstration of a synthetic pathway that can support photosynthesis in two ways. First, it can produce acetyl-CoA from C<sub>3</sub> sugars without releasing CO<sub>2</sub>. It can also assimilate photorespiratory glycolate without loss of carbon (Yu *et al.*, 2018).

### **Carbon fixation via a novel pathway *in vitro***

An *in vitro* reconstruction of a synthetic carbon fixing pathway, the CETCH cycle, based on highly efficient reductive carboxylation. The pathway, utilizing 17 enzymes that originate from 9 organisms, was optimized by a combination of enzyme engineering and metabolic proofreading (Schwander *et al.*, 2016).

### **Overexpressing the H-protein of the glycine cleavage system increases biomass yield in glasshouse and field grown transgenic tobacco plants**

Increased biomass upon overexpression of a limiting photorespiratory protein in tobacco grown in field conditions. This indicates that optimization of expression levels within native carbon fixation-related pathways could be harnessed to increase productivity, and that photorespiration could be improved even without the need for synthetic pathways (Lopez-Calcano *et al.*, 2018).

### **The road to C<sub>4</sub> photosynthesis: evolution of a complex trait via intermediary states**

A case for engineering C<sub>3</sub>-C<sub>4</sub> intermediate metabolism as a way to increase photosynthetic efficiency and set the stage towards future realization of complete C<sub>4</sub> metabolism. This study suggests that a detailed and mechanistic understanding of C<sub>3</sub>-C<sub>4</sub> intermediates could provide valuable guidance for experimental designs aiming to boost carbon fixation (Schlüter and Weber, 2016).

### **Evolving *Methanococcoides burtonii* archaeal Rubisco for improved photosynthesis and plant growth**

A demonstration of the use of directed laboratory evolution to improve the kinetic properties of Rubisco from an archaeal origin. The improved Rubisco variant was introduced to tobacco chloroplast and demonstrated to increase photosynthesis. Such protein engineering strategies could be used to address the kinetic limitations of key enzymes, thus supporting higher metabolic fluxes and boosting productivities (Wilson *et al.*, 2016).

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