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- 17
- 18 One-sentence summary
- 19 Photosynthetic efficiency of soybean pods and seeds contribute to the seed weight and the
- 20 overall carbon gain of the canopy by compensating for carbon loss, with implications for yield.

21 ABSTRACT

There is a limited understanding of the carbon assimilation capacity of non-foliar green 22 tissues and its impact on yield and seed quality since most photosynthesis research focuses on 23 24 leaf photosynthesis. In this study, we investigate the photosynthetic efficiency of soybean (*Glycine max*) pods and seeds in a field setting and evaluate its effect on mature seed weight and 25 26 composition. We demonstrate that soybean pod and seed photosynthesis contributes 13-14% of the mature seed weight. Carbon assimilation by soybean pod and seed photosynthesis can 27 28 compensate for 81% of carbon loss through the respiration of the same tissues, and our model predicts that soybean pod and seed photosynthesis contributes up to 9% of the total daily carbon 29 30 gain of the canopy. Chlorophyll fluorescence shows that the operating efficiency of Photosystem II in immature soybean seeds peaks at the 10-100mg seed weight stage, while that of immature 31 32 pods peaks at the 75-100mg stage. This study provides quantitative information about the efficiency of soybean pod and seed photosynthesis during tissue development and its impact on 33 yield. 34

35

36 INTRODUCTION

Photosynthesis converts light energy to transform CO_2 into soluble carbohydrates, which 37 are then utilized for plant growth and maintenance (Stirbet et al., 2020). Most research has 38 focused on leaf photosynthesis with a minimal understanding of potential carbon assimilation in 39 40 non-foliar green tissue and its contribution to yield and seed quality. Chlorophyll-containing seeds (Schwender et al., 2004), ears of wheat (Triticum aestivum) (Maydup et al., 2010), and 41 husks of maize (Zea mays) (Pengelly et al., 2011), along with other green non-foliar tissues, such 42 as stems, all perform photosynthesis, possibly providing an additional important source of 43 44 photoassimilates for the plant. Non-foliar photosynthesis is a photosynthetic process similar to that of the foliar mesophyll cell (Simkin et al., 2020), except that there are potentially two 45 46 important sources of CO₂ for non-foliar photosynthesis. Both foliar and non-foliar photosynthesis utilize ribulose-1,5-bisphosphate carboxylase/oxygenase (Rubisco), which assimilates 47 48 atmospheric CO₂ that enters the cells through the stomata. In contrast to foliar tissue, some nonfoliar tissues fix a larger portion of CO_2 that is released by mitochondrial respiration and then 49

refixed by Rubisco (Aschan and Pfanz, 2003; Millar et al., 2011). Stomata are found in varying numbers on non-foliar tissues; however, their function and the amount of photosynthesis that relies on CO_2 from the atmosphere through these stomata have yet to be determined (Simkin et al., 2020).

The embryos of many plants (e.g., Arabidopsis (Arabidopsis thaliana), chickpea (Cicer 54 55 arietinum), coffee (Coffea arabica), cotton (Gossypium hirsutum), soybean (Glycine max), oilseed rape (Brassica napus), pea (Pisum sativum), broad bean (Vicia faba)) contain 56 57 considerable amounts of chlorophyll, although the level of chlorophyll depends on the developmental stage (Yakovlev and Zhukova, 1980; Simkin et al., 2010; Puthur et al., 2013; 58 59 Smolikova and Medvedev, 2016). Green embryos, known as chloroembryos (Palanisamy and Vivekanandan, 1986), have all the photosynthetic complexes such as Photosystem I (PSI), 60 61 Photosystem II (PSII), cytochrome b6f complex, and ATP synthase (Weber et al., 2005; Allorent et al., 2015; Kohzuma et al., 2017). Rubisco is known to be active in seeds of soybean (Allen et 62 al., 2009), oilseed rape (Hills, 2004; Ruuska et al., 2004), broad bean, and fenugreek (Trigonella 63 foenum-graecum) (Willmer and Johnston, 1976). Seed photosynthesis contributes to the 64 accumulation of storage lipids in oilseed rape (Eastmond et al., 1996; Ruuska et al., 2004), and 65 66 the rapid production of ATP and NADPH for the synthesis of complex carbohydrates, fatty acids, and proteins has also been linked to photosynthesis in some chloroembryos (Asokanthan et 67 al., 1997; Wu et al., 2014). 68

69 The tissue surrounding a seed, such as a pod in legumes, may also photosynthesize, and 70 the rate depends heavily on the species and how much light its seed-related tissues receive, 71 which is determined by plant and canopy architecture. Rubisco activity has been detected in the 72 pod walls of peas; however, Rubisco concentrations are 10-100 times lower than that seen in leaf tissue (Hedley et al., 1975). The photosynthetic activity of the wheat ear substantially contributes 73 74 to the pool of carbohydrates translocated to the developing grains during the post-anthesis stages 75 (Tambussi et al., 2005; Tambussi et al., 2007; Maydup et al., 2010; Sanchez-Bragado et al., 2014). Although the wheat ear net CO₂ assimilation rate is lower on an area basis than that of the 76 flag leaf (Tambussi et al., 2005; Tambussi et al., 2007), wheat ear photosynthesis can contribute 77 up to 70% of the individual grain weight yield component in a wide range of genotypes (Maydup 78 et al., 2010) and contrasting environments (Sanchez-Bragado et al., 2014). Shading research in 79

barley (*Hordeum vulgare*) demonstrates a large contribution of the ear to grain weight (up to
50%) and thus yield, similar to wheat (Bort et al., 1994). Despite that only a fraction of the above
canopy photosynthetic photon flux density (PPFD) reaches legume embryos, seed coats, and
pods, all three tissues have relatively high rates of electron transport (Allen et al., 2009;
Tschiersch et al., 2011). Both immature soybean pods and seeds are green and capable of
photosynthesis but unlikely to experience positive net rates of photosynthesis (Sambo et al.,
1977) due to high rates of respiration.

87 The most frequent method for determining the contribution of non-foliar photosynthesis is the suppression of photosynthesis through light exclusion (i.e., covering) or herbicide 88 89 treatment (Sanchez-Bragado et al., 2016). Seeds that develop in foil-wrapped Arabidopsis siliques have lower oil and starch, as well as lower overall seed weight, than seeds that develop 90 91 in siliques exposed to light (Liu et al., 2017). Other Arabidopsis studies using chemical inhibitors of embryonic photosynthesis (Allorent et al., 2015) or inducible photosynthesis-deficient mutants 92 93 (Sela et al., 2020) report that final seed lipid and protein content is unaffected but that germination and early growth from the seeds are reduced. 94

Here, we investigate the photosynthetic capacity of soybean pods and seeds during
development and evaluate the impact of pod and seed photosynthesis on seed weight and
composition under field conditions, covering developing soybean pods with aluminum foil for 68 weeks (starting from less than 2 cm in pod length to fully grown and desiccated). Our goal is to
determine whether pod and seed photosynthesis in soybean contributes significantly to the
mature seed and to the plant overall.

101

102 **RESULTS**

Pod and seed photosynthesis contributes 13-14% of the seed weight under Illinois field conditions

To understand the effect of pod and seed photosynthesis on the seed weight and
 composition of soybean under field conditions, we conducted field experiments on covered pods.
 We covered pods of Clark (maturity group IV) soybean with foil at an early stage of

108 development to block light from reaching the pods and seeds and then measured the weight and

- 109 composition of the mature soybean seeds that developed in those pods compared to uncovered
- 110 pods (Figure 1A and B). The results for the cultivar Clark showed that covering the pods with
- 111 foil during development significantly reduced the final weight of the seeds inside (14.3 g)
- 112 compared to seeds from uncovered pods (16.6 g), a decrease of 13% (Figure 1C). Similar results
- 113 were found with another cultivar in the field, Williams 82 (maturity group III), in which seeds
- 114 from foil-covered pods showed a significant reduction in final seed weight (14.5 g) compared to
- seeds from uncovered pods (16.9 g), a decrease of 14% (Figure 2B).

116 We compared the efficiency of photosystem II between immature covered and uncovered pods and seeds of Williams 82 from the field using chlorophyll fluorescence. We collected green 117 pods and seeds at 100-200 mg (average fresh seed weight). The foil-covered pods and seeds were 118 119 visibly much paler (Supplemental Figure S1A and D), confirmed by the chlorophyll fluorescence images (Supplemental Figure S1B and E). The operating efficiency of photosystem II (F_q '/ F_m ') 120 was significantly reduced in both the seeds and the pods that had developed in foil, compared to 121 the uncovered seeds and pods (Supplemental Figure S1C and F). The temperature and humidity 122 123 of the air space inside the foil cover and immediately outside were measured and found to have 124 no significant difference (Supplemental Figure S2), indicating temperature and humidity did not affect the change in seed weight. 125

The effect of covering on the concentrations of protein and oil was not consistent in the two genetic backgrounds. Seeds from covered pods had a 5% decrease in oil and a 5% increase in protein compared to seeds from uncovered pods in Clark (Figure 1D and E), but no significant difference was found in Williams 82 (Figure 2C and D). No seed composition parameter evaluated here (Supplemental Figures S3 and S4) appeared consistently affected by the foilcovering across the cultivars tested.

132

Pod and seed photosynthesis can compensate for 81% of carbon loss from respiration of pod and seed under field conditions

We measured rates of net photosynthesis and dark respiration (R_{dark}) in pods during development (Figure 3A) using the LI-6800 (LICOR Biosciences Inc, Lincoln, NE, USA), allowing illumination on both sides of the pod (Figure 3B). For the determination of gross photosynthesis, the rate of respiration in the light was assumed to be equal to the rate of respiration in the dark. Measured rates of dark respiration were significantly higher in uncovered pods (10.7 µmol CO₂ m⁻² s⁻¹) compared to covered pods (8.3 µmol CO₂ m⁻² s⁻¹) (Table 1).

We estimated the rate of gross photosynthesis, the true rate of photosynthesis, taking into 141 account the rate of respiration, by adding R_{dark} to the rate of net photosynthesis at each PPFD 142 143 (Wittmann et al., 2006). Rates of net photosynthesis were low in illuminated pods that developed 144 in the dark for 3-4 weeks and were substantially different across PPFD levels, following a pattern typical for light-response curves in leaves (Figure 3C). The highest measured rates of net and 145 gross photosynthesis occurred at 2500 μ mol quanta m⁻² s⁻¹ for uncovered and 750 μ mol quanta 146 m⁻² s⁻¹ for covered pods (Figure 3D). Measured net photosynthesis in uncovered pods peaked at a 147 rate of 0.48 μ mol CO₂ m⁻² s⁻¹, whereas measured net photosynthesis in covered pods peaked at a 148 rate of -5.9 μ mol CO₂ m⁻² s⁻¹. In uncovered pods, measured rates of gross photosynthesis peaked 149 at 11.2 μ mol CO₂ m⁻² s⁻¹, and covered pods peaked at a rate of 2.4 μ mol CO₂ m⁻² s⁻¹. Rates of 150 saturating photosynthesis were modeled using the R package 'Photosynthesis' 151 (Stinziano et al., 2020); the modeled rates of saturating gross photosynthesis were similar to 152 measured rates of gross photosynthesis. Modeled saturating rates of gross photosynthesis were 153 11.4 μ mol CO₂ m⁻² s⁻¹ and 2.5 μ mol CO₂ m⁻² s⁻¹ in uncovered and covered pods, respectively 154 (Table 1). The maximum quantum efficiency (ΦCO_2), estimated from modeling, was ten times 155 higher in uncovered pods than in covered pods (Table 1). 156

157 Canopy incident light for this experiment was considered to be 300 µmol quanta m⁻² s⁻¹ 158 (close to 260 µmol quanta m⁻² s⁻¹ measured by Allen et al., 2009) for uncovered and 0 µmol 159 quanta m⁻² s⁻¹ for covered pods. The rates of net photosynthesis at canopy incident light were 160 measured as -2 µmol CO₂ m⁻² s⁻¹ for uncovered and -8.3 µmol CO₂ m⁻² s⁻¹ for covered pods 161 (Table 1). The rates of gross photosynthesis in these pods at canopy incident light levels were 162 measured as 8.7 µmol CO₂ m⁻² s⁻¹ and 0 µmol CO₂ m⁻² s⁻¹ (Table 1) for uncovered and covered, 163 respectively. While uncovered pods at the canopy light condition of 300 µmol quanta m⁻² s⁻¹ were still a net carbon loss for the plant, pod and seed photosynthesis was nevertheless able to compensate for 81% of the carbon lost through respiration.

166

A canopy model projects that pod and seed photosynthesis contributes up to 9% of total gross canopy photosynthesis

169 Soybean pods are mostly located under the canopy, where various light environments are 170 present depending on height, developmental stage and time of day. To estimate the contribution 171 of pod photosynthesis to overall canopy photosynthesis of soybean, model simulation studies were performed using a multilayer canopy model (Campbell & Norman, 1998; Drewry et al., 172 173 2010). The measured leaf area index was 9.52 ± 0.20 (SE), and the pod area index was $1.30 \pm$ 174 0.14 (SE); the leaves were mainly concentrated in the middle of the canopy, while the pods were more evenly distributed throughout the canopy (Figure 4A, Supplemental Table S1). Simulation 175 176 of canopy photosynthesis over the whole day of 17 August 2022 (R6 stage, full-length filled 177 pods and fully green plant) indicated that pod photosynthesis contributed about 9.6% to the overall gross assimilation of CO₂ of the soybean canopy (Figure 4B). The simulated proportion 178 of pod contribution was greater in the early morning and late afternoon and slightly less at noon 179 180 (8.1%).

181

Efficiency of seed and pod photosynthesis peaks by the 100 mg developmental stage under field conditions

We investigated the photosynthetic efficiency of immature seeds from field-grown 184 Williams 82 soybeans during their development (Figure 5A). We collected immature pods and 185 measured seed fresh weight to separate seeds into seven different developmental stages (Figure 186 5B), then measured the operating efficiency of photosystem II (F_a'/F_m') of the seeds (Figure 5C). 187 The operating efficiency of photosystem II (F_q'/F_m') of the seeds started at 0.323 in the 5-10 mg 188 stage, increased in the 10-25 mg stage (0.393), and remained steady before decreasing (0.363) in 189 the 75-100 mg developmental stage (Figure 5D). The chlorophyll concentration of seeds started 190 191 at 0.426 mg/g DW in the 5-10 mg stage, decreased in the 10-25 mg stage (0.287 mg/g DW), then

increased and remained steady until the 75-100 mg stage (0.432 mg/g DW), after which it

decreased (Figure 5E). The ratio of chlorophyll a to b increased across the sequential

194 developmental stages (Figure 5F).

195 We also investigated the photosynthetic efficiency of the corresponding developing pods by measuring the efficiency of photosystem II in the same way after the seeds had been removed. 196 197 The developmental stages of pods were labeled based on the average weight of seeds inside the pods (Figure 6B). The operating efficiency of photosystem II of the pods started at 0.291 in the 198 199 5-10 mg stage, decreased in the 10-25 mg stage (0.273), then rose to a peak at the 75-100 mg stage (0.3) (Figure 6D). The concentration of chlorophyll in these pods started at 0.63 mg/g DW 200 201 in the 5-10 mg stage, then gradually decreased until the 75-100 mg stage (0.355 mg/g DW), after which it rose slightly (Figure 6E). The ratio of chlorophyll a to b remained similar across the 202 203 sequential pod developmental stages (Figure 6F).

204

Photosynthesis-related genes were actively expressed in the early stages of seed development

We explored which photosynthesis genes were expressed in early soybean seed 207 development using previously published high-throughput transcriptome profiling (RNA-seq) 208 with the cultivar Williams 43 (Jones et al., 2013; Cho et al., 2019). The expression levels of 209 photosynthesis-related genes were disproportionately high in the very young seeds, mostly over 210 211 50 RPKM (Reads Per Kilobase per Million mapped reads), whereas most genes in general (>98%) were expressed lower than 50 RPKM in the data. The gene expression levels were 212 mostly high in the early stages (a few days after flowering) and peaked at 5-6 mg in whole seeds, 213 including genes related to Rubisco, the light reactions, the photosynthetic carbon metabolism 214 215 reactions, and chlorophyll synthesis (Figure 7, Supplemental Figures S5 and S6). At later stages (100-200 mg and older), photosynthesis-related genes were actively expressed but mostly 216 217 decreased in the cotyledon tissues.

Interestingly, the gene expressions of photosynthesis-related genes (Figure 7) were active ahead of the peak photosynthetic efficiency of the seeds at the 10-100 mg stages, as shown by

chlorophyll fluorescence (Figure 5C). Expression levels of genes related to chlorophyll cycle 220 pathways peaked before and after the 5-6 mg stage (Supplemental Figure S6C). This occurrence 221 222 may explain the decrease of chlorophyll levels at the 10-25 mg stage (Figure 5E). Chlorophyllrelated genes were expressed highly in different developmental stages depending on their 223 functions. Chlorophyll synthesis-related genes were highly expressed in the early stages and then 224 225 decreased after 5-6 mg, while chlorophyll degradation and heme synthesis-related genes were active at the latest developmental stages when the seed had turned yellow and was drying down 226 (Supplemental Figure S6). These data suggest that the efficiency of photosynthesis and the level 227 of chlorophyll are tightly connected with gene expression during seed development. 228

229

230 **DISCUSSION**

In this study, we evaluated the effect of soybean pod and seed photosynthesis on seed 231 232 weight and composition in a field setting and also investigated the photosynthetic efficiency and 233 rate during seed development. This study demonstrated that soybean pod and seed photosynthesis contributes 13-14% of the seed weight in the field in Illinois, which is a major 234 soybean growing area (Figures 1-2). The carbon assimilation by pod and seed photosynthesis at 235 236 under-canopy light levels can compensate for 81% of carbon loss through the respiration of the 237 same tissue (Figure 3). A multilayer canopy model predicts that pod and seed photosynthesis contributes up to 9% of the daily total carbon assimilation of the soybean canopy (Figure 4). 238 Chlorophyll fluorescence showed that the operating efficiency of PSII in immature seeds peaked 239 at the 10-100 mg seed weight stages, while that of immature pods peaked at the 75-100 mg seed 240 weight stage (Figure 6). Previously, there was little understanding of potential carbon 241 assimilation in seed and pod tissues and its impact on yield and seed quality. 242

243 Despite having low rates of net photosynthesis, uncovered soybean pods have the ability 244 to fix more CO₂ than is lost through dark respiration at light levels greater than 245 1000 μ mol quanta m⁻² s⁻¹. However, under canopy light conditions at 300 μ mol quanta m⁻² s⁻¹, 246 the rate of net photosynthesis in uncovered pods was measured at -2 μ mol CO₂ m⁻² s⁻¹, and gross 247 photosynthesis was measured at 8.7 μ mol CO₂ m⁻² s⁻¹ (Figure 3 and Table 1). While uncovered 248 pods under canopy light conditions were still a net carbon loss for the plant, our data showed that pod photosynthesis was able to compensate for 81% of the carbon lost through respiration under normal field within canopy light levels. The ΦCO_2 of pods in Williams (0.06) is similar to the ΦCO_2 of leaves in Clark (0.051-0.068, from Slattery et al. 2017), indicating that pods and leaves have a similar quantum efficiency for carbon assimilation. In contrast, the gross photosynthesis at saturating light in pods was about 3 times lower than that in the leaves (11.4 vs 35.7 µmol CO₂ m⁻² s⁻¹). The capacity of photosynthesis on an area basis is much higher in leaves, while the quantum efficiency of carbon assimilation is similar between the two tissues.

Covering young soybean pods with foil and allowing the seeds to develop to maturity in 256 the foil, thereby reducing the input from photosynthesis, resulted in seeds with a lower weight 257 (g/100 seeds) than uncovered seeds by 13-14%. This effect was seen in two different cultivars of 258 soy that were grown in the Illinois field (Figures 1 and 2). From this, we conclude that seed and 259 pod photosynthesis contributes to soybean seed weight under field conditions, similar to results 260 found for wheat ears (Maydup et al., 2010) and barley ears (Bort et al., 1994). The relative 261 importance of these non-foliar assimilates varies depending on the species, cultivar, and 262 environment (Tambussi et al., 2007). From our data, we conclude that green soybean pods and 263 264 seeds are photoheterotrophic, meaning that seeds and pods are sink tissues, but also that their photosynthesis substantially contributes to carbon gain, being responsible for 9% of total daily 265 266 seed carbon according to our model.

Our model estimated the contribution of pod and seed photosynthesis from the 267 268 parameters of a maturity group III variety measured at the SoyFACE field in Illinois on 17 August 2022 (93 days after planting), when the deepest canopy was present during the 269 development of soybean. The angle of the sun during early morning and late afternoon allows a 270 greater proportion of light to reach the pods (Supplemental Table S2), which in turn leads to a 271 272 higher proportion of pod contribution during those times (Figure 5). Pods under the canopy can get more light before or after the development of the densest canopy; thus, our model likely 273 274 tends to underestimate the contribution of pod and seed photosynthesis to canopy photosynthesis. We set the pod angle distribution parameter in the multilayer canopy model at zero, thus 275 276 assuming all pods are vertical for convenience, which is another factor by which our model may 277 underestimate the contribution of pod and seed photosynthesis because the angle of pods in the field is greater than zero. Our model was able to parameterize the sun leaf angle allowing better 278

light penetration into the canopy in the morning and afternoon, leading to the proportion of pod
contribution being greater in the early morning and late afternoon than at noon. Threedimensional (3D) canopy photosynthesis models can accurately simulate the intricate 3D
architecture and microclimate under the canopy during plant development (Watanabe et al.,

283 2005; Zheng et al., 2008; Song et al., 2013; Wang et al., 2017; Wang et al., 2020).

284 We cannot exclude the possibility that limiting light from the developing pods and seeds may have impacts not directly related to photosynthetic carbon gain. One possibility is that the 285 286 foil covering interferes with the development of plastids, which play a critical role in seed development as a source of energy for oil synthesis and starch storage. Despite the low levels of 287 available light inside the seed, soybean seed photosynthetic activity is surprisingly high, possibly 288 due to the unique nature of the chloroplasts present in the seed (Borisjuk and Rolletschek, 2009). 289 290 These chloroplasts have chlorophyll-protein complexes similar to those found in leaf chloroplasts but with a higher proportion of granal stacking (Saito et al., 1989; Asokanthan et al., 291 292 1997), which allows for distinct light harvesting properties and low saturation levels for photosynthetic electron transport (Borisjuk et al., 2005). The seed chloroplasts conduct light 293 294 reactions to generate ATP/NADPH, which can fuel the biosynthesis of seed oil (Ruuska et al., 295 2004; Schwender et al., 2004). Furthermore, a recent study showed that the chloroplasts of bean fruit mesocarp cells differentiate into amyloplasts, which allows for the storage of surplus 296 sucrose for starch synthesis, later used by the seeds when the sucrose supply is no longer 297 298 sufficient (Belmont et al., 2022).

299 Recent studies using different methods to block pod and seed photosynthesis in 300 Arabidopsis have found that mature seed lipid and protein are not affected, whether the 301 photosynthesis is blocked by chemicals (Allorent et al. 2015) or by inducible mutations (Sela et al. 2020). In contrast, Liu et al. (2017), who wrapped developing Arabidopsis siliques in foil, 302 303 observed a decrease in oil bodies in the dark-grown embryos, and in other species, such as B. 304 napus, reduced pod and seed photosynthesis has a negative impact on the amount of seed oil (Ruuska et al. 2004; Schwender et al., 2004). We evaluated the effect of pod and seed 305 photosynthesis on soybean seed composition however could not conclude that light levels and 306 related pod and seed photosynthesis affect soybean seed composition (Figures 1 and 2). 307 Numerous studies have reported that soybean seeds at the top of the canopy contain more protein 308

and less oil than seeds at the bottom of the canopy (Collins & Carter, 1956; Escalante & Wilcox, 309 1993a and 1993b; Huber et al., 2016). We collected seeds from the four uppermost nodes 310 311 (designated as the upper layer of the canopy) separately from the rest of the nodes (designated as the lower layer of the canopy). The results for both of the cultivars studied here (Clark and 312 Williams 82) showed that covering the pods with foil during development significantly reduced 313 314 the final seed weight, compared to seeds from uncovered pods wherever they were located in the canopy (Supplemental Figures S7 and S8). In contrast to the seed weight, the protein and oil 315 316 concentrations were unaffected by the foil-covering and instead related only to the position of the pod in the canopy. However, as we did not track the position (upper or lower) of the foil-covered 317 pods, we cannot say for certain that the foil-covering had no effect on seed composition. 318

This study investigated the photosynthetic efficiency of developing soybean pods and 319 320 seeds by measuring $F_{q'}/F_{m'}$, representing the operating efficiency of PSII photochemistry, which is indicative of functional electron transport. Modulated chlorophyll fluorescence assesses the 321 photosynthetic efficiency indirectly but more easily and quickly than other methods, such as gas 322 exchange, and has been previously used for this purpose in wheat ear studies (Tambussi et al., 323 324 2005; Maydup et al., 2012). Our results showed that seed photosynthesis peaked at the 10-100 325 mg developmental stages and that pod photosynthesis peaked at the 75-100 mg developmental stage in Illinois field conditions (Figures 5 and 6). Additionally, the expression levels of 326 photosynthesis-related genes were high in the early stages of seed development, even earlier than 327 were studied here with chlorophyll fluorescence, and peaked at the 5-6 mg developmental stage, 328 329 then decreased with time (Figure 7). The expression of chlorophyll synthesis genes at older seed stages, such as 100-200 mg, was observed to be synchronized with the fluorescence 330 measurements and chlorophyll amounts in this study (Figures 5-7). These observations suggest, 331 332 unsurprisingly, that the efficiency of photosynthesis and the production of chlorophyll are tightly connected with gene expression during seed development, similar to leaf photosynthesis (Foyer 333 et al., 2012; Hibberd and Covshoff, 2010). 334

Multinational efforts have been made to improve photosynthesis over the past decade, with the premise that crop production improvements must be swift and substantial due to the plateauing of yields of important crops in the face of rapidly growing agricultural demand (Long and Ort, 2010). Several studies reported improved photosynthesis and increased yield in model

plants (Kromdjik et al., 2016; South et al., 2019) and crops (Yoon et al., 2020; de Souza et al., 339 340 2022). One way to improve photosynthesis in crops is to engineer a better light distribution under the canopy so that photosynthetic tissues receive more light and can operate at higher efficiency 341 (Ort et al., 2015). Improved light distribution inside a canopy could also increase the light use 342 efficiency of non-foliar tissues such as soybean pods and seeds. Although leaves are the most 343 important photosynthetic tissue, this study and others (Simkin et al., 2020; Lawson and Miliken, 344 2023) suggest that improved efficiency of non-foliar tissues, such as soybean pods and seeds, can 345 346 also contribute to improved yield of important crops.

In summary, we demonstrated that pod and seed photosynthesis contributed a significant portion of carbon assimilation and seed weight in soybean–9% of total daily carbon assimilation according to our model and 13-14% of seed weight, which is a vital component of yield. These data suggest that pod and seed photosynthesis plays an important role in soybean yield, which merits further study in the context of improving global food supplies in a sustainable way.

352

353 MATERIALS AND METHODS

354 Field conditions

Two cultivars of soybean [*Glycine max* (Williams 82 PI518671 and Clark PI548533)] were planted in 1.5-m rows using standard agronomic practices at the University of Illinois Energy Farm field station (40.11°N, 88.21°W, Urbana, IL, USA) on 27 May 2021. Each row (1 block) consisted of ~10 plants spaced 3.8 cm apart. There were 4 blocks for Clark and 8 blocks for Williams 82.

360

361 **Treatment and sample collection**

A cluster of young pods (smaller than 2 cm, the earliest in their development that we were able to wrap with foil without risk of damage) was carefully wrapped in aluminum foil with the intent to block all light from reaching the pods, while still allowing them to grow. Multiple clusters of pods were covered with foil per block in the field, while leaving the majority of the

pods on the plant uncovered. Pods remained covered in foil for at least 3 weeks before collection 366 367 (for immature tissues) or until maturity (6-8 weeks). Immature pods were collected and opened, and the seeds were removed. The immature seeds were weighed to determine the developmental 368 stage (fresh weight range); pods were labeled by the fresh weight of the seeds inside them (using 369 the stage of the majority of the seeds if not all were the same). Foil-covered seeds and pods were 370 371 kept in the dark as much as possible during the procedure. For the Williams 82 field-grown developmental series, immature seeds (and their pods) were collected at 7 fresh weight ranges: 5-372 10 mg, 10-25 mg, 25-50 mg, 50-75 mg, 75-100 mg, 100-200 mg, 200-300 mg. Dry pods were 373 harvested at maturity and shelled, and the seeds were further dried in the oven overnight at 50 °C 374 or air-dried for at least 2 weeks before further measurements. 375

376

Mature seed harvest and non-destructive near-infrared (NIR) spectroscopy for seed composition

379 From the field, both Clark and Williams 82 pods were harvested at maturity on 4 October 2021. Uncovered and foil-covered pods were collected separately. Each block for each genotype 380 contained approximately 10 plants, and seeds were pooled from plants within the same 381 382 block/genotype/treatment. Uncovered pods were initially divided into 'Up' (harvested from the upper four nodes of the plant) and 'Low' (harvested from the rest of the plant). For Clark, seeds 383 from foil-covered pods were obtained from all 4 blocks (one block was removed from data 384 analysis as an outlier); for Williams 82, seeds from foil-covered pods were obtained from 5 of 8 385 blocks. Uncovered and foil-covered seeds were further dried in the oven overnight at 50 °C, then 386 air-dried for at least 2 weeks before NIR measurement. For seeds from uncovered pods, total 387 388 weight and the weight of 100 randomly-selected seeds were obtained from each block (divided into 'Up' and 'Low'); from this, an approximate seed count was calculated. For seeds from foil-389 covered pods, the exact number of seeds per block was counted and weighed. 390

Near-infrared (NIR) spectroscopy was performed using the Perten DA7250
(PerkinElmer, Waltham, MA, USA) with the company settings for whole soybean seeds, nondestructively measuring 28 components including protein, oil, 5 fatty acids, and 18 amino acids.
The mirror cup was used with all samples. For the Clark and Williams 82 field-grown seeds from

uncovered pods, 20 replicates of 10 seeds each were measured per block (200 different seeds per block), half from the 'Up' seeds and half from the 'Low' seeds. 'Up' and 'Low' data were later combined for analysis. For the Clark seeds from foil-covered pods, 4 blocks yielded 1 to 3 replicates each, for a total of 7 replicates; each replicate had 10 seeds. For the Williams 82 seeds from foil-covered pods, 5 blocks yielded 1 to 5 replicates each, for a total of 11 replicates; each replicate had 10 seeds.

401

402 **Pod photosynthesis measurements**

Photosynthesis on pods still attached to the plants was measured between 7 September 403 404 and 15 September 2021 at the UI Energy Farm. Pods were measured using an LI-6800 (LICOR 405 Biosciences Inc, Lincoln, NE, USA) fitted with a clear-top chamber (LI-6800-12A) using the 3 cm X 3 cm aperture insert. Small light sources (LI-6800-02) were fitted to the top and bottom of 406 407 the clear-top chamber to allow illumination on both sides of the pod (Fig 3A). Pods were placed flat in the chamber to allow for full illumination on both sides of the pod. Pods were 408 photographed, and projected pod area was estimated using ImageJ (US National Institutes of 409 Health, Bethesda, MD, USA). The pod was allowed to acclimate in the chamber for at least 15 410 411 minutes or until the rate of photosynthesis was stable. Reference CO_2 was set to 400 µmol mol⁻¹, PPFD set to 2500 μ mol quanta m⁻² s⁻¹, energy balance set to 30°C, and relative humidity set to 412 50%. The chamber was set to an overpressure of 0.1 kPa with a flow rate of 1100 μ mol s⁻¹. Rates 413 of net photosynthesis were measured across a light-response curve which consisted of the 414 following PPFDs for both light sources: 2500, 2000, 1600, 1200, 900, 750, 600, 500, 400, 300, 415 200, 120, 60, 20, 0, 0, 0, 0 μ mol quanta m⁻² s⁻¹ with a minimum wait time of 60 s and a 416 maximum wait time of 90 s that the LI-6800 was set to match before each measurement. We 417 assumed the rate of respiration in the light was equal to the rate of dark respiration (R_{dark}). Gross 418 photosynthesis, the true rate of photosynthesis, was estimated by adding R_{dark} to the rate of net 419 photosynthesis at each PPFD (Wittmann et al., 2006). Transmittance through pods is higher than 420 421 transmittance through leaves (Allen et al., 2009); however, the transmittance through the entire seed is low. Most of the area taken up by the pod in the chamber included the seeds, leading us to 422

423 choose to present PPFD on a single-sided basis as light is unlikely to pass through the seed.

424 Light-response curves were measured on five covered and five uncovered Williams 82 pods.

425

426 Leaf area index (LAI) measurements

Leaf area index was measured on 17 August 2022 using the Sunscan Canopy Analysis System (DeltaT Devices, Cambridge, UK) at 12 cm height increments within the soybean canopy. Six rows of soybean were measured, where four replicate positions were averaged for each increment. The Ellipsoidal Leaf Angle Distribution Parameter (ELADP) was set to 0.81, and absorption was set to 0.85.

432

433 Canopy model

To estimate the contribution of pod photosynthesis to the overall canopy photosynthesis of soybean, model simulation studies were performed using a multilayer canopy model (Campbell & Norman, 1998; Drewry et al., 2010). Leaf area indexes were measured from the SoyFACE field in Illinois on 17 August 2022. Pod area indexes were estimated by measuring plant density, pod area, and number of pods per plant (Supplemental Table S1). The leaf angle distribution parameter was set as 0.81 (Campbell & Norman, 1998), and the pod angle distribution parameter was set as 0, assuming all pods are vertical.

441

442 Chlorophyll fluorescence for photosystem II operating efficiency

443 F_q'/F_m' was measured by a chlorophyll fluorescence (CF) imager (CF Imager, 444 Technologica, UK). Pods (with seeds removed) were exposed to 260 µmol quanta m⁻² s⁻¹ (light 445 intensity under canopy from Allen et al., 2009) for 10 min until stabilized, then subjected to a 446 burst of 6100 µmol quanta m⁻² s⁻¹ for 800 ms. Seeds were exposed to 78 µmol quanta m⁻² s⁻¹ 447 (light intensity inside pods from Allen et al., 2009) for 10 min until stabilized then subjected to a 448 burst of 6100 µmol quanta m⁻² s⁻¹ for 800 ms. After the burst, *F*' and *F_m*' were automatically

measured by the software. The images were manually adjusted, and F_q'/F_m' was calculated. For 449 the Williams 82 field-grown developmental series, each of the 7 immature seed stages included 450 451 41 to 48 individual seeds, and each of the 7 immature pod stages included 32 to 42 pod halves across 7 blocks (1 block was not used). Seeds were removed from the pods prior to imaging. 452 One-half of the pod was imaged, interior side down. Seeds and pods were collected and imaged 453 over two days, 25 to 26 August 2021. Immature Williams 82 field-grown foil-covered pods were 454 collected and imaged over two days, 8 and 10 September 2021. Four immature foil-covered pods 455 (7 pod-halves) and their 12 seeds (100-200 mg fresh weight) were imaged separately. For 456 comparison, 5 immature uncovered pods (7 pod-halves) and their 15 seeds (100-200 mg fresh 457 weight) collected at the same time were imaged separately. 458

459

460 Chlorophyll measurement

461 Separated immature pod and seed samples were frozen in liquid nitrogen for at least 10 minutes and stored in the freezer (-80 °C) until they were lyophilized. Chlorophyll content was 462 determined using 100% (v/v) ethanol extraction (Ritchie, 2006) and microplate spectrometer 463 (Warren et al., 2008). For the Williams 82 field-grown developmental series, each of the 7 464 465 immature seed stages included 4 to 10 replicates (containing 3 to 4 individual seeds each) from 2 to 4 (of 8) different blocks. Each of the 7 immature pod stages included 6 to 12 replicates (an 466 individual pod-half as a replicate) from 3 to 6 (of 8) different blocks. Lyophilized seeds were 467 weighed, and weight was considered when calculating the amount of chlorophyll present. 468

469

470 Transcriptome data analysis

All raw high-throughput transcriptome (RNA-Seq) data were downloaded from the Gene Expression Omnibus (GEO, GSE42871 and GSE123655); see Supplemental Dataset S1 for the list of samples. Photosynthesis-related soybean gene models were chosen based on a keyword search of annotations at Phytozome (https://phytozome-next.jgi.doe.gov/).

476 **Temperature and humidity measurements**

Temperature and humidity were measured inside 17 foil-covered packets surrounding
pods in the field using SRH77A Temperature / Humidity Thermistor Instrument (Cooper-Atkins,
CT, USA). An equal number of measurements were taken from equivalent pods that were
uncovered. Field measurements were taken on 10 September 2021.

481

482 Statistical analysis

The statistical analyses were done with R (version 4.0.3). The biomass, photosynthesis 483 estimated from fluorescence, and chlorophyll levels from field trials were analyzed in a mixed 484 model analysis of variance (ANOVA) followed by a *post-hoc* Tukey test (α =0.05). Block was 485 considered as the random effect for all analyses, while foil-covering treatment was considered as 486 the fixed effect. For the light-response curves, we used the lme4 R package (Bates et al., 2015) to 487 perform linear mixed effects analyses of the relationship between the physiological response 488 489 variables (net and gross photosynthesis) and treatment (covered vs. uncovered). We set treatment 490 and PPFD as fixed effects. We structured the model to allow for random intercepts for individual 491 pods. Rates of saturating gross photosynthesis, dark respiration and ρCO_2 were modeled using the 'photosynthesis' R package (Stinziano et al. 2020). Saturating rates of net and gross 492 photosynthesis and measured rates of dark respiration were compared in covered and uncovered 493 pods using independent t-tests. If data were not normally distributed, non-parametric analysis 494 (Wilcoxon rank test) was conducted by R program (α =0.05). 495

496 Data availability

- 497 Raw data available in the Supplemental Dataset S1 File.
- 498 Supplemental Data
- 499 Supplemental Figure S1. The efficiency of photosystem II of covered and uncovered pods
 500 under field conditions (Williams 82).
- 501 **Supplemental Figure S2.** Temperature and humidity around covered and uncovered pods.

Supplemental Figure S3. Seed composition of covered and uncovered seeds under field
 conditions (Clark).

Supplemental Figure S4. Seed composition of covered and uncovered (lower or upper position)
 seeds under field conditions (Wm82).

- Supplemental Figure S5. Expression of genes annotated as Rubisco, Photosystem I,
 Photosystem II, light harvesting complex, cytochrome b6 f-related, fructose bisphophatase and -
- 508 aldolase (Calvin cycle), ferredoxin-related, ATPase.
- 509 Supplemental Figure S6. Expression of genes annotated as chlorophyll synthesis.
- 510 Supplemental Figure S7. Seed composition of covered and uncovered seeds under field511 conditions (Clark).
- 512 Supplemental Figure S8. Seed composition of covered and uncovered seeds under field
 513 conditions (Wm82).
- Supplemental Table S1. Values, units and descriptions of the parameters in the multilayer
 canopy model of soybean.
- 516 **Supplemental Table S2.** The total light absorbed by leaves and pods of the soybean canopy.
- 517 Supplemental Dataset S1.

519

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- 535 Y.W., D.R.O. wrote the paper.

536

537 **TABLES**

Table 1. Pod photosynthetic parameters for Williams 82, measured 7 September through 16 September 2021 at the University of Illinois Energy Farm in Urbana, IL, USA. Different letters represent significant differences at the p<0.05 level obtained from independent t-tests. Asterisk (*) indicates model parameters

541 were estimated using the R package 'Photosynthesis' (Stinziano et al., 2020).

	Uncovered	Covered
Respiration in the dark (umol CO ₂ m ⁻² s ⁻¹)	10.7 ^a	8.3 ^b
Gross photosynthesis at light saturation*	11.4 ^a	2.5 ^b
(μmol CO ₂ m ⁻² s ⁻¹) ΦCO ₂ gross photosynthesis at light acturation*	0.06 ^a	0.007 ^b
Net photosynthesis at incident	-2	-8.3
light (μ mol CO ₂ m ⁻² s ⁻¹)	(at 300 μ mol quanta m ⁻² s ⁻¹)	(at 0 μ mol quanta m ⁻² s ⁻¹)
Gross photosynthesis at incident	8.7	0
light (µmol CO ₂ m ⁻² s ⁻¹)	(at 300 μ mol quanta m ⁻² s ⁻¹)	(at 0 μ mol quanta m ⁻² s ⁻¹)

545

546 FIGURE LEGENDS

547 Figure 1. Seed composition of covered and uncovered seeds under field conditions (Clark). The box plots

show the median (central line), the lower and upper quartiles (box), and the minimum and maximum

549 values (whiskers). Each dot represents a value (n= 3 blocks). N.s., not significantly different (α =0.05).

550 ANOVA is used with linear mixed model (random effect = block, fixed effect = cover). Least squares

551 mean is used to compare. Covering affects seed weight, oil and protein percentage.

552 A. Picture of soybean in the field (Clark).

B. Representation of aluminum foil that was used to cover pods, when the length of pod was shorter than2 cm.

555 C. Seed weight, g/100 seeds. Seeds from covered pods: $14.3g \pm 0.4$ (SE). Seeds from uncovered pods:

556 $16.6g \pm 0.4$ (SE). Seeds from covered pods have a 13% decrease in weight compared to seeds from 557 uncovered pods.

- 558 D. Oil, percent. Seeds from covered pods: 18.7%. Seeds from uncovered pods: 19.8%. Seeds from
- 559 covered pods have a 5% decrease in oil compared to seeds from uncovered pods.
- E. Protein, percent. Seeds from covered pods: 46.1%. Seeds from uncovered pods: 43.7%. Seeds from
- 561 covered pods have an 5% increase in protein compared to seeds from uncovered pods
- 562
- 563 **Figure 2.** Seed composition of covered and uncovered seeds under field conditions (Wm82). The box
- plots show the median (central line), the lower and upper quartiles (box), and the minimum and maximum
- values (whiskers). Each dot represents a value (n= 5 blocks for covered and 8 blocks for uncovered).
- 566 ANOVA is used with linear mixed model (random effect = block, fixed effect = cover, α =0.05). N.s., not
- significantly different (α =0.05). Least squares mean is used to compare. Covering affects seed weight but

568 not oil or protein percentage.

569 A. Picture of soybean at the field (Wm82).

- 570 B. Seed weight, g/100 seeds. Seeds from covered pods: $14.5g \pm 0.6$ (SE). Seeds from uncovered pods:
- 571 $16.9g \pm 0.4$ (SE). Seeds from covered pods have a 14% decrease in weight compared to seeds from 572 uncovered node
- 572 uncovered pods.
- 573 C. Oil, percent. Seeds from covered pods: 20.5%. Seeds from uncovered pods: 20.9%. There is no
- 574 significant difference in oil between the seeds from the covered and the uncovered pods.
- 575 D. Protein, percent. Seeds from covered pods: 43.2%. Seeds from uncovered pods: 43.8%. There is no
- 576 significant difference in protein between the seeds from the covered and uncovered pods.
- 577
- 578 **Figure 3**. Rates of net photosynthesis and dark respiration in pods. Small leaf and needle chamber with
- 579 the small light sources to allow illumination of the pod from the top and bottom (A). Pods were placed
- 580 horizontally in the chamber to allow full illumination and to estimate pod area (B). Rates of net (C) and
- 581 gross (D) photosynthesis of covered (white) and uncovered (black) pods under field conditions (Williams
- 582 82). Each dot represents a value $(n=5) \pm SE$. We assumed that the seeds greatly inhibited the transmittance
- 583 of light through the pod and used photosynthetic photon flux density (PPFD) for a single-side.

584

585 **Figure 4**. Canopy model of soybean photosynthesis. A. Leaf area index and pot area index at each layer

- of canopy from SoyFACE soybean field. Layer1 is the top and level 8 is the bottom. B. Predicted diurnal
 gross CO2 assimilation of leaf and pod in the soy canopy on 17 August 2022.
- 588

589 **Figure 5**. Photosynthetic activity of seeds in seven developmental stages under field conditions: 5-10 mg,

590 10-25 mg, 25-50 mg, 50-75 mg, 75-100 mg, 100-200 mg, and 200-300 mg, based on the fresh weight of 591 the seeds.

- 592 A. Picture of soybean (Williams 82) plant at the 2021 Illinois field.
- 593 B. Picture of collected soybean seeds with fresh weight range shown to the right. 5-10 mg seeds are on
- 594 the top row with successively larger seeds in the lower rows.
- 595 C. Chlorophyll fluorescence (CF) image of photosystem II operating efficiency (Fq'/Fm') of seeds. This
- 596 is the same plate shown in B.
- 597 D. Average value of Fq'/Fm' with error bars representing standard error (n=41-48).
- 598 E. Average value of total chlorophyll level with error bars representing standard error (n=4-10).
- 599 F. Average value of chlorophyll a/b ratio with error bars representing standard error (n=4-10).
- 600
- 601 **Figure 6**. Photosynthetic activity of pods in seven developmental stages under field conditions: 5-10 mg,
- 602 10-25 mg, 25-50 mg, 50-75 mg, 75-100 mg, 100-200 mg, 200-300 mg. Cultivar is Williams 82.
- 603 A. Picture of soybean plant at the 2021 Illinois field.
- 604 B. Picture of collected soybean pods. The fresh weight range of the seeds inside the pods is shown to the 605 left and right.
- 606 C. Chlorophyll fluorescence (CF) image of photosystem II operating efficiency (Fq'/Fm') of pods. This is
- 607 the same plate shown in B.

- D. Average value of Fq'/Fm' with error bars representing standard error (n=32-42).
- 609 E. Average value of total chlorophyll level with error bars representing standard error (n=6-12).

Figure 7. Expression of photosynthesis related genes during seed development. A-B. Rubisco genes (dark

610 F. Average value of chlorophyll a/b ratio with error bars representing standard error (n=6-12).

611

613 614	reaction); C-D. Light harvesting complex genes (light reaction); E-F. Mg-chelatase genes (chlorophyll synthesis); G-H. fructose bisphophatase and -aldolase (Calvin cycle) genes. Developmental stages for whole good (WS): 1 - 4 Days After Fertilization (DAF) WS: 2 - 12 - 14 DAF WS: 2 - 22 - 24 DAF WS: 4 - 5
015 616	whole seed (WS): 1=4 Days After Fertilization (DAF) WS; 2=12-14 DAF WS; 3=22-24 DAF WS; 4=5- 6mg WS, Developmental stages for cetyleden: 4=5.6mg cetyleden: 5=100.200mg cetyleden: 6=400
617	500mg cotyledon: 7–200-300mg vellow cotyledon: 8–dry cotyledon. Beads Per Kilobase per Million
618	mapped reads (RPKM).
619	
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