

1 **Title:** Whole plant chamber to examine sensitivity of cereal gas exchange to changes  
2 in evaporative demand

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4 **Authors:** Iván Jauregui, Shane A Rothwell, Samuel H Taylor, Martin AJ Parry,  
5 Elizabete Carmo-Silva, Ian C Dodd

6

7 **Address:** Lancaster Environment Centre, Lancaster University, Lancaster, LA1 4YQ,

8 UK

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16 **Corresponding author:**

17 Name: Ivan Jauregui

18 Address: Lancaster Environment Centre, Lancaster University, Lancaster, LA1

19 4YQ, United Kingdom

20 Current address: Genetic lab. Gembloux Agro Bio-Tech, University of Lieje,

21 Gembloux, 5030, Belgium

22 E-mail address: [i.jauregui@lancaster.ac.uk](mailto:i.jauregui@lancaster.ac.uk); add here new email too

23 Name: Ian Dodd

24 Address: Lancaster Environment Centre, Lancaster University, Lancaster, LA1

25 4YQ, United Kingdom

26 E-mail address: [i.dodd@lancaster.ac.uk](mailto:i.dodd@lancaster.ac.uk)

27 **ABSTRACT**

28           Background: Improving plant water use efficiency (WUE) is a major target for  
29 improving crop yield resilience to adverse climate change. Identifying genetic variation  
30 in WUE usually relies on instantaneous measurements of photosynthesis ( $A_n$ ) and  
31 transpiration ( $T_r$ ), or integrative measurements of carbon isotope discrimination, at the  
32 leaf level. However, leaf gas exchange measurements alone do not adequately  
33 represent whole plant responses, especially if evaporative demand around the plant  
34 changes.

35           Results: Here we describe a whole plant gas exchange system that can rapidly  
36 alter evaporative demand when measuring  $A_n$ ,  $T_r$  and intrinsic WUE (iWUE) and  
37 identify genetic variation in this response.  $A_n$  was not limited by VPD under steady-  
38 state conditions but some wheat cultivars restricted  $T_r$  under high evaporative  
39 demand, thereby improving iWUE. These changes may be ABA-dependent, since the  
40 barley ABA-deficient mutant (*Az34*) failed to restrict  $T_r$  under high evaporative  
41 demand. Despite higher  $T_r$ , *Az34* showed lower  $A_n$  than wild-type (WT) barley  
42 because of limitations in Rubisco carboxylation activity.  $T_r$  and  $A_n$  of *Az34* were more  
43 sensitive than WT barley to exogenous spraying with ABA, which restricted  
44 photosynthesis via substrate limitation and decreasing Rubisco activation.

45           Conclusions: Examining whole plant gas exchange responses to altered VPD  
46 can identify genetic variation in whole plant iWUE, and facilitate an understanding of  
47 the underlying mechanism(s).

48

## BACKGROUND

49           Photosynthesis is a complex process in which light, water and carbon dioxide  
50 (CO<sub>2</sub>) are used to synthesize carbohydrates. In plants, CO<sub>2</sub> can only diffuse into the  
51 leaves via the stomata. When open, the stomata represent the major path of water  
52 loss to the atmosphere via transpiration. Approximately 98% of all water taken up  
53 through the roots may be transpired through the stomata [1]. Therefore, plants  
54 constantly seek to minimise water loss while maintaining CO<sub>2</sub> entry for photosynthesis,  
55 by tightly regulating their stomatal responses. Monitoring plant–atmosphere gas  
56 exchange is essential for understanding plant responses to a fluctuating environment.

57           Atmospheric vapour pressure deficit (VPD) or evaporative demand is  
58 influenced by both air temperature and relative humidity (RH), and is the difference  
59 between the saturation vapour pressure and the actual vapour pressure. The driving  
60 force for water movement through the plant is caused by the vapour pressure deficit  
61 between the substomatal cavity and the surrounding air, known as leaf-to-air vapour  
62 pressure deficit (VPD<sub>leaf</sub>). High VPD<sub>leaf</sub> increases plant transpiration rates (Tr) [2]. By  
63 decreasing their stomatal conductance (gs), plants can partially limit Tr and the  
64 decrease in leaf water status [3]. High ambient VPD and VPD<sub>leaf</sub> enhances evaporation  
65 of water from the leaf, reducing bulk leaf water status and inducing stomatal closure,  
66 which is contributed to by a hydropassive response common to all land plants and, in  
67 angiosperms, a hydroactive response regulated by abscisic acid (ABA) [4]. Increased  
68 VPD rapidly upregulates expression of the *NCED* genes (involved in ABA  
69 biosynthesis), thereby increasing leaf [ABA] and decreasing gs [5]. However, this leaf-  
70 based mechanism may not completely explain the spatial and temporal behaviour of  
71 whole plant transpiration under increasing evaporative demand: other factors such as  
72 patchy stomatal closure [6], changes in leaf [7], root [8,9], or whole plant hydraulic

73 conductivity [10,11] and leaf-age differences in sensitivity to ABA [12], may operate  
74 together to limit  $T_r$  under increasing VPD.

75 Water use efficiency (WUE) typically refers to the ratio between the biomass  
76 produced and cumulative water use. At the physiological level, the ratio of net  
77 photosynthesis ( $A_n$ ) to  $T_r$  is known as photosynthetic or intrinsic WUE (iWUE).  
78 Maintaining net photosynthesis ( $A_n$ ) while reducing  $T_r$  under high atmospheric  
79 evaporative demand may be of adaptive significance under certain conditions, and  
80 genetic variability in the sensitivity of  $g_s$  to VPD has been described in angiosperms:  
81 in some genotypes,  $T_r$  increases linearly with increasing VPD, while others restrict  $T_r$   
82 at higher VPD. Pioneering work identified the “restricted transpiration” trait [13, 14],  
83 and associated low leaf hydraulic conductivity with improved WUE. The trait has been  
84 identified in many crops, including cereals [15–16], using gravimetric methods in  
85 chambers [17], greenhouses [18], and the field [19]. A potential drawback of  
86 decreasing  $g_s$  to restrict transpiration under increasing VPD, is that internal  $CO_2$   
87 concentration ( $C_i$ ) may decrease, thereby decreasing  $A_n$  via substrate limitation. Field  
88 measurements under high VPDs cannot separate effects of VPD on  $A_n$  from effects  
89 of high temperature *per se*. Consistent with this potential limitation, high evaporative  
90 demands and temperatures considerably limit leaf level photosynthesis [20,21].  
91 However, similar measurements at the whole plant level have not been made.

92 Leaf gas exchange measurements fail to capture whole plant responses since:  
93 <sup>1)</sup> transpiration inside the leaf cuvette of an infra-red gas analysis system reflects the  
94 controls imposed on that environment (i.e. mixing of air to control boundary layer  
95 conductance, chosen temperature, choice of light source, leaf area used for  
96 measurement, flow rate); <sup>2)</sup> leaf measurements cannot adequately describe whole  
97 plant  $A_n$  due to spatial variation in the light environment of different leaves [22,23]; <sup>3)</sup>

98 naturally occurring microclimates across the plant affect its interaction with the  
99 environment. Thus, several chambers have been built to characterize whole plant gas  
100 exchange of plants such as Arabidopsis [24–26], shrubs [27–29], or even trees [30],  
101 but with limited regulation of environmental conditions inside the chamber. As a  
102 consequence, such measurements may be bedeviled by leaks, flow rate fluctuations,  
103 overheating of the larger chambers [31], and high humidity/condensation that can  
104 cause severe failures of IRGAs [32,33]. These technical difficulties probably explain  
105 why relatively few researchers have built whole plant systems to study transpiration  
106 responses to increasing evaporative demand [7,18,34,35].

107         In the present manuscript, we describe a whole plant gas exchange system to  
108 measure  $A_n$ ,  $T_r$  and  $iWUE$  under increasing VPD. We tested whether different cereal  
109 genotypes, previously demonstrated to show variation in transpiration response to  
110 VPD [16] and variation in leaf-level photosynthesis [36], showed variation in whole  
111 plant  $iWUE$  as evaporative demand changed. Because higher photosynthetic rates  
112 correlate with high yield [36] and stomatal responses to VPD governs diurnal plant  
113 transpiration [39], identifying useful genetic variation in  $iWUE$  at high VPD will be of  
114 interest to plant physiologists and breeders. Our whole plant gas exchange system is  
115 relevant to achieving this goal.

116         Since the role of ABA in regulating stomatal responses to VPD is not completely  
117 clear (cf. [35,37]), we used the whole plant gas exchange system to investigate the  
118 responses of an ABA-deficient barley mutant and its wild-type under contrasting VPD,  
119 and in response to foliar ABA spraying. Previous observations indicate that exogenous  
120 ABA application limits photosynthesis of ABA-deficient plants (*flacca* tomato mutant  
121 and Arabidopsis lines) [35, 38], even if the mechanistic interpretation is not clear. Our  
122 working hypothesis is that stomatal hypersensitivity of the ABA deficient mutant (*Az34*)

123 to exogenous ABA spraying constrains photosynthesis via substrate limitation,  
124 decreasing Rubisco activation state, and limiting net photosynthesis.

125

## 126 **MATERIA METHODS**

### 127 **Growth conditions and plant material**

128 Wheat (*Triticum aestivum*) and barley (*Hordeum vulgare*) were pre-germinated  
129 on moistened filter paper (Whatman #1) in petri dishes. The dishes were covered with  
130 foil and placed in dark conditions at room temperature ( $24^{\circ}\text{C} \pm 5\%$ ) for 48 h. Once  
131 germinated, two seeds were placed at about 2.5 cm depth in rectangular 2 l pots (10.5  
132 x 10.5 x 20 cm height) containing a commercial growing substrate (Petersfield  
133 Products, UK) with a slow-release fertilizer (Osmocote, Scotts UK Professional, UK).  
134 After the first true leaf emerged, one of the plants was removed from each pot to  
135 maintain one plant per pot. Twelve days after transplanting, the plants were supported  
136 in a sealing sleeve (Fig. 1, Supplemental Fig. 1). The plants grew for six weeks until  
137 reaching the phenological stage Zadoks 39-45. Plants were watered every 2-3 days  
138 to reach drip point, the maximum water content of the substrate, and were randomly  
139 allocated in the greenhouse and rotated weekly to assure homogeneity.

140 Plants were placed in a naturally lit greenhouse at Lancaster University  
141 ( $54.0104^{\circ}\text{N}$ ,  $2.7877^{\circ}\text{W}$ ) with supplementary lighting (14 h per day), and controlled  
142 temperature (lights turn off if air temperature exceeds  $30^{\circ}\text{C}$ ). To maintain atmospheric  
143 VPD lower than 2.5 kPa throughout a diurnal cycle, a ten heads humidifier (Growell,  
144 UK) was placed in the greenhouse, to avoid developmental VPD priming of plants  
145 growing in different periods in the greenhouse.

146 Table 1 describes the different experiments done. The wheat cultivars (cv.)  
147 Krichauff and Drysdale were chosen because they showed contrasting  $T_r$  under  
148 increasing VPD [16]. The wheat cultivars cv. Cadenza, Gatsby, Mercato, Gladiator,  
149 Zebedee were chosen because they showed contrasting leaf photosynthesis (A) in a  
150 field experiment [36]. The barley ABA-deficient mutant *Az34* mutant (and its  
151 corresponding wild-type, WT) was chosen since it shows reduced capacity to produce  
152 ABA in response to water deficit, caused by a pleiotropic deficiency in the molybdenum  
153 cofactor that decreases aldehyde oxidase activity, which catalyses the ultimate step  
154 in the ABA biosynthesis pathway [40]. This mutant has higher  $T_r$  than the wild-type  
155 (WT) Steptoe in an early stage, both under control VPD and after increasing air  
156 temperature and, therefore, VPD [41].

157

### 158 **Whole plant gas exchange system**

159 We re-designed the whole plant gas exchange system previously described  
160 [18]. With the new configuration and upgrades, the equipment can measure  $A_n$  and  
161  $iWUE$ , in addition to  $T_r$ , under increasing VPD (Fig. 1). Hereafter, transpiration  
162 determined with this chamber is termed  $T_{IRGA}$  to avoid confusion with  $T_r$  obtained by  
163 gravimetric methods. The new system incorporates: <sup>1)</sup> a powerful  
164 humidifier/dehumidifier system (Supplemental Fig. 5) that can more rapidly change  
165 chamber relative humidity (5 min compared to ~30 min required previously [18])  
166 allowing higher VPDs (> 4 kPa) while maintaining temperatures below 30°C; <sup>2)</sup> a mass  
167 flow controller to tightly control the flow in the system by allowing a certain amount of  
168 pressure from the compressor while the previous version [18] pulled in air via a fan <sup>3)</sup>  
169 multiple probes within the chamber to monitor environmental conditions including



170 temperature, relative humidity and light, which were absent in [18]; <sup>4</sup>) a LI- 6400XT (LI-  
171 COR, Lincoln, NE, USA) to simultaneously measure the gas exchange by logging the  
172 data measured using the various probes. The diagrams of the different parts are  
173 supplied in Supplemental Fig.s 1-4.

174

### 175 **IRGA and external probes**

176 A LI-6400XT equipped with a 9964-053 Sample Cell Outlet Manifold Kit (LI-  
177 COR, Lincoln, NE, USA) to reduce the gas analyzer sample volume, was used to  
178 determine CO<sub>2</sub> and H<sub>2</sub>O vapour concentrations. Using a LI-6400XT allowed external  
179 probes to be connected to the console to calculate A<sub>n</sub> and T<sub>IRGA</sub>, using a protocol  
180 provided by LI-COR (LI-6400 Portable Photosynthesis System, Application Note 2) to  
181 communicate with the external probes as well as the IRGA. A temperature-humidity  
182 probe (Vaisala Humitter 50Y, Helsinki, Finland), a flow rate transducer (TSI 8450,  
183 Aliflow Instruments, USA) and a temperature probe (LI-COR, Lincoln, Nebraska, USA)  
184 were added.

185 The LI-6400XT head was connected to the chamber using vinyl flexible tubing  
186 (Swagelok, UK) and aluminium tube fittings and adapters (Swagelok, UK). Gas was  
187 driven through the LI-6400XT head using an external pump (model TD-4X2NA,  
188 Brailsford & CO, USA), which tightly controlled the flow of air. The flow rate achieved  
189 was checked every week. All tubing was covered with thermal insulation to stabilise  
190 dew point temperatures.

191

### 192 **The chamber**

193 A chamber of total volume of 30 l (25 x 20 x 60 cm) was built from Perspex,  
194 with a nominal thickness of 3.5 mm. Light was supplied by two Son-T high-pressure  
195 sodium lamps (Philips, Netherlands) providing  $450 \mu\text{mol m}^{-2} \text{s}^{-1}$  PPFD at the top of  
196 the plants. The light action spectrum that it is transmitted into the chamber was  
197 measured between 200 nm and 1100 nm by placing the spectroradiometer (SR9910-  
198 V7, Macam Photometrics, Livingston, UK) inside the closed chamber, at 25 nm  
199 intervals (Supplemental Fig. 6).

200 To insert the plant into the chamber, one side consisted of a removable door  
201 (see Supplemental Fig. 3a) sealed with 1 cm wide neoprene sponge rubber and  
202 closed using eight metal clips. A sealable slot at the base of the chamber (see sealing  
203 sleeve description) isolated the root and shoot of the plant.

204 The chamber is hermetically sealed and works under a slight overpressure. It  
205 is noteworthy that no leaks were detected in our system (Fig. 2A-B). Four fans  
206 (Ebmpapas 512Ft, Hungary) were placed inside the chamber (two fans on the top  
207 quarter and two fans on the bottom quarter of the chamber) to lower boundary layer  
208 resistance, with a combined capacity of  $310 \text{ m}^3 \text{ h}^{-1}$ . Fan placement ensured  
209 homogenous airflow which was checked using smoke (data not shown). The  
210 equipment is operated in the laboratory, allowing the temperature to remain stable at  
211  $27.5^\circ\text{C} \pm 5\%$  when the fans are on (Fig. 2C). Temperature and relative humidity inside  
212 the chamber remains comparable (Supplemental Table 1).

213

#### 214 **Sealing sleeve**

215 A sealing sleeve, made of PVC (12 x 8 x 0.2 cm) (Supplemental Fig. 1) isolated  
216 the above and below-ground parts of the plant. In most cases, tiller development inside

217 the sealing sleeve isolated the roots from the shoots, but to ensure gas tightness  
218 Sylgard Silicone elastomer (Dow Corning, UK) was applied inside the sealing sleeve  
219 two days prior to measurements. A neoprene sponge rubber ensured a tight fit of the  
220 plant into the chamber.

221

## 222 **The circuit**

223 Air from outside the building was supplied to the chamber to assure a stable  
224 [CO<sub>2</sub>]. The [CO<sub>2</sub>] in this source changed less than 10 ppm during a typical day. To  
225 provide air under positive pressure, we used a compressor (OF1202-40MQ3, Junk Air,  
226 USA) with an extensive cooling system for temperature control. The compressed air  
227 was circulated through a 2 m pipe (1 cm internal  $\varnothing$ ) filled with silica gel to dehydrate it  
228 to approx. 5% RH. The silica gel was replenished after every 4-6 h of use. Thus  
229 conditioned, the air was supplied to the chamber at a stable rate of 30 l min<sup>-1</sup> (with an  
230 error of 0.1%) via a mass flow controller (Alicat CMR 500 SLPM, Alicat Scientific,  
231 USA). This flow rate allows <sup>1)</sup> an acceptable air renewal of one chamber volume per  
232 minute; <sup>2)</sup> a reasonable [CO<sub>2</sub>] differential across the chamber (between -18 to -25  $\mu\text{mol}$   
233 CO<sub>2</sub> m<sup>-2</sup> s<sup>-1</sup>); <sup>3)</sup> avoidance of high system pressure.

234 If necessary (at high VPDs or when plant leaf area exceeded 400 cm<sup>2</sup>), the flow  
235 rate was increased. A water bath (Fig. 1, Supplemental Fig. 2) containing an Ultrasonic  
236 humidifier with ten heads (Growell, UK) was developed to re-humidify the air (if  
237 needed). Manually operated low-pressure valves (Swagelok, UK) were used to control  
238 the amount of air passing through the water bath. The system established RH values  
239 in the range 5-75% by passing air through the water bath, and when higher RH was  
240 desired, the ultrasonic humidifier was connected. Although most of the tubing in the

241 system has 0.4 cm internal  $\varnothing$ , the tube that connects the humidifier system with the  
242 pre-mixer chamber has 1 cm internal  $\varnothing$  to avoid condensation inside it.

243 To homogenize the air, it passes through a pre-mixer box (30 x 30 x 30 cm)  
244 (Fig. 1, Supplemental Fig. 4). Next, prior to entering the chamber, the air transits a  
245 PVC pipe (3 cm internal  $\varnothing$ , 40 cm length) where flow rate, temperature and humidity  
246 probes and the reference line for LI-6400XT are assembled. The flow rate used for the  
247 gas exchange calculations is computed here. Typically, the conductance of this pipe  
248 averages  $175 \mu\text{mol air s}^{-1}$  with an error of 5%. The pipe ends in the base of the  
249 chamber, circulating air upwards. Air exits the chamber via another PVC pipe where  
250 the sample line for the LI-6400XT is connected. A thermocouple (connected to the LI-  
251 6400XT) measures the temperature of a selected leaf from the top of the canopy.

252

### 253 **Data collection**

254 At the beginning of each measurement sequence, the plants were acclimated  
255 for  $\sim 20$  min to a VPD of 2.5 kPa, the maximum VPD experienced by plants in the  
256 greenhouse. Differences in  $[\text{CO}_2]$  and  $[\text{H}_2\text{O}]$  between air entering and exiting the  
257 chamber were measured and recorded using the LI-6400XT. Once the exchange of  
258  $\text{CO}_2$  and  $\text{H}_2\text{O}$  had been steady for more than 5 min (steady-state, Fig. 2D), values  
259 were logged every 20 seconds for 3-5 min, and a median value was established. Then,  
260 the relative humidity in the system was adjusted to inside the system were changed to  
261 achieve the next desired VPD level, usually requiring 15-30 min to reach a new steady-  
262 state. For VPD curves, VPD was gradually decreased to the minimum achievable in  
263 decrements. After that, VPD was increased in 0.5 kPa (or 0.75 kPa) increments to a  
264 maximum of 3.75 kPa during winter and above 4 kPa during summer experiments.

265 Each plant was exposed to a minimum of 7 different VPDs. After measuring whole  
266 shoot gas exchange response to changing VPD, each plant was removed from the  
267 chamber to determine leaf area (LI-3100C Area Meter, Lincoln, NE, USA).  $T_{IRGA}$  did  
268 not significantly differ from gravimetrically determined  $T_r$  (see Supplemental Fig. 7)

269 To examine the effects of the *Az34* mutation in barley plants, leaf [ABA] was  
270 measured as previously described [12]. Frozen leaf tissues were freeze-dried and  
271 then powdered in a mortar. The ABA was extracted in distilled water (1:50, w/w) at 4  
272 °C overnight in a shaker. ABA concentration was determined in aqueous extracts by  
273 a radioimmunoassay with the monoclonal antibody MAC252 as previously described  
274 [42]. The assay was conducted with two technical replicates per biological sample  
275 (Supplemental table 2).

276 In some experiments, ABA was sprayed on the leaves to inhibit  $T_r$ . ABA was  
277 dissolved in ethanol to make a stock solution at 0.05 M, which was diluted to 10  $\mu$ M in  
278 H<sub>2</sub>O prior to use. ABA 10  $\mu$ M was applied with a wetting agent Silwet (L-77, De  
279 Sangosse Ltd, Cambridge, UK) at 0.025 %. We applied 10-15 ml per plant, depending  
280 on leaf area, using an atomizer (Perfume Pod, Amazon, UK). ABA-sprayed plants  
281 were used to measure whole plant gas exchange after 1 h (Supplemental Fig. 7).

282 Flag leaf gas exchange measurements were also made as part of these  
283 experiments spraying ABA over whole plants. The conditions in the LI-6400XT  
284 chamber were 1.5 kPa air VPD (to avoid stomatal limitations at high VPD), 500  $\mu$ mol  
285 s<sup>-1</sup> air flow, 400 ppm CO<sub>2</sub>, 25°C leaf temperature (same as the *in vitro* Rubisco assay)  
286 and 460  $\mu$ mol m<sup>-2</sup> s<sup>-1</sup> PPFD.

287 Flag leaf Rubisco *in vitro* activity was measured with a non-radioactive  
288 spectrophotometric assay with the modifications described by [43, 44]. The assay uses

289 five enzymatic reactions to couple ribulose 1,5-bisphosphate (RuBP) carboxylation  
290 and 3-PGA formation to NADH oxidation. Rubisco activity is calculated based on  
291 RuBP consumption by monitoring the decrease in NADH concentration in the well,  
292 tracking the absorbance at 340 nm using UV-transparent 96-well plates in a microplate  
293 reader (SpectroStars, BMG Labtech, Germany) at 25°C. Firstly, leaves were extracted  
294 as described by [36]. The Rubisco total activity ( $V_t$ ) was assayed after incubating the  
295 extract for 5 min in the presence of CO<sub>2</sub> and MgCl<sub>2</sub>, while the initial activity ( $V_i$ ) was  
296 measured directly after extraction. The Rubisco activation state is the ratio  $V_i / V_t$ .

297

## 298 **Statistical analysis**

299 One- or two-way ANOVA [45] was used to test statistical significance of  
300 differences in means of each trait between genotypes or between genotypes and ABA  
301 treatments, respectively. Where significance of effects was observed ( $P < 0.05$ ),  
302 multiple pairwise comparisons between treatments used the Tukey-b test.

303 To detect the  $T_{IRGA}$  breakpoint, the R package “segmented” [46] was used.  
304 When the results lacked biological meaning (resulting from statistical artefacts  
305 associated with exceeding the VPD operating boundaries of the chamber), or when  
306 the slope after the breakpoint was higher than the one before, a linear regression was  
307 used. Breakpoint calculations were made for each plant individually (Supplemental  
308 tables 3-5). Regression results were confirmed using the software Prism 7 (GraphPad  
309 Software Inc., San Diego, USA; Supplemental table 6).

310

## 311 **RESULTS**

## 312 **Reliability of the whole plant system**

313 Without a plant inside the chamber,  $\Delta H_2O$  and  $\Delta CO_2$  were stable over  
314 time at a steady flow rate (Fig. 2A), and at different flow rates (Fig. 2B), indicating  
315 that leaks were absent or minimal. Rapid and large VPD changes (0.5-4 kPa) were  
316 possible in just a few min (Supplemental Fig. 5) while maintaining a temperature of  
317  $27.5^\circ C \pm 5\%$ , which is faster than in previously reported chambers [7,18,34,35].

318 The whole plant system shows similar stability with a plant inside the chamber.  
319 Temperature and relative humidity were stable over time (Fig. 2C) because <sup>1)</sup> the  
320 system was mounted in a temperature-controlled laboratory, and <sup>2)</sup> the pipes were  
321 thermally insulated. Moreover, the water bath design (Supplemental Fig. 2) assured  
322 stability of VPD (Fig. 2C). With a wheat plant in the chamber, whole plant  $A_n$  and  
323  $T_{IRGA}$  remained stable over time (Fig. 2D).  $T_{IRGA}$  measurements did not produce  
324 different results from paired gravimetric measurements (Supplemental Figure 7).

325

## 326 **Whole plant gas exchange at a single VPD**

327 At a single VPD ( $2.5 \text{ kPa} \pm 0.15$ ) and constant temperature ( $27.5^\circ C \pm 1\%$ ),  
328 different wheat cultivars showed significant differences in whole plant gas exchange  
329 (Fig. 3). Transpiration varied ca. 17%, with Cadenza having higher  $T_{IRGA}$  than  
330 Mercato, Zebedee and Gladiator, while Gatsby, Drysdale and Krichauff, had  
331 intermediate values (Fig. 3A). Photosynthesis varied ca. 30%, with Gatsby having  
332 higher  $A_n$  than Gladiator, Krichauff and Cadenza, while Mercato, Zebedee and  
333 Drysdale had intermediate values (Fig. 3B).  $iWUE$  was more influenced by whole plant  
334  $A_n$  than  $T_{IRGA}$  (Fig. 3C). Gatsby and Zebedee had higher  $iWUE$  than Cadenza,  
335 Gladiator and Krichauff, with Mercato and Drysdale having intermediate values. Since

336 whole plant iWUE of Drysdale and Krichauff was similar, their gas exchange was  
337 studied under contrasting VPD levels.

338

### 339 **Effects of changing VPD on whole plant gas exchange**

340 A representative example of the data required to examine the presence of the  
341  $T_{IRGA}$  breakpoint (BP) is shown in Fig. 4. Measurements commenced at 2.5 kPa, the  
342 VPD experienced by plants in the greenhouse; then VPD was decreased to the  
343 minimum achievable in 0.5 kPa steps (Fig. 4A). After that, VPD was increased in 0.5-  
344 0.75 kPa steps to a maximum of 3.75 kPa. Air temperature inside the chamber  
345 remained stable during data collection (Fig. 4A). Following this protocol, plant gas  
346 exchange usually equilibrates within about 15-30 min because of the small (usually  
347 0.5 kPa) VPD changes over time, and because, as grasses, *Triticum spp.* show  
348 relatively rapid stomatal movement due to their stomatal conformation [47]. Each VPD  
349 response curve took 3-4 h, and no pronounced hysteresis was detected when plants  
350 were exposed to ascending and descending series of VPDs (Supplemental Fig. 9).  
351 To avoid hydraulic limitations of transpiration that occur if the upper layers of the  
352 substrate dry out [48], water was added to the pot every hour during measurement  
353 until leaching was observed (since the pot could be irrigated without opening the  
354 chamber).

355

### 356 **Restriction of whole plant gas exchange under high vapour pressure deficit**

357 In the wheat cv. Drysdale,  $T_{IRGA}$  increased with increasing VPD and showed a  
358 BP at  $2 \pm 0.3$  kPa ( $R^2 = 0.96$ ), while in cv. Krichauff,  $T_{IRGA}$  increased linearly with VPD  
359 ( $R^2 = 0.91$ ) (Fig. 5A,D; Supplemental Table 3), as previously described [16]. Across



360 the entire range of VPDs,  $T_{IRGA}$  was significantly ( $P = 0.002$ ) higher in cv. Krichauff  
361 than cv. Drysdale. When comparing  $T_{IRGA}$  below the Drysdale BP, both cultivars  
362 showed similar sensitivity of  $T_{IRGA}$  to VPD (same slope,  $P = 0.21$ ). Beyond this BP,  
363 the slopes significantly differ ( $P = 0.003$ ) with  $T_{IRGA}$  less sensitive to VPD in cv.  
364 Drysdale. Drysdale plants had a significantly ( $P = 0.003$ ) higher An. Taken together,  
365 cv. Drysdale had a significantly ( $P = 0.005$ ) higher iWUE than cv. Krichauff over the  
366 entire VPD range.

367 In wild-type (WT) barley, transpiration increased linearly with VPD up to  $1.9 \pm$   
368  $0.3$  kPa ( $R^2 = 0.96$ ), but VPDs above this threshold restricted transpiration (Fig. 5G,  
369 Supplemental Table 3). In contrast, transpiration of the ABA-deficient *Az34* barley  
370 mutant increased linearly and continuously with increasing VPD ( $R^2 = 0.91$ ) (Fig. 5J).  
371 Absolute  $T_{IRGA}$  of the *Az34* mutant was similar to WT over the entire VPD range.  
372 Before the BP at 1.9 kPa, the slope of the  $T_{IRGA}$  versus VPD response was similar ( $P$   
373  $= 0.55$ ) between genotypes; beyond this BP,  $T_{IRGA}$  was more sensitive to VPD in the  
374 WT ( $P = 0.0037$ ). An was significantly lower in *Az34* than WT plants ( $P < 0.001$ ), and  
375 decreased as VPD increased for the mutant only (Fig. 5 H,K; Supplemental Table 4).  
376 *Az34* plants had a significantly ( $P = 0.002$ ) lower An than WT both before ( $P < 0.001$ )  
377 and after ( $P < 0.001$ ) the BP (1.9 kPa). In *Az34* plants, iWUE decreased exponentially  
378 as VPD increased, while in WT plants, iWUE decreased as VPD increased in the low  
379 range, but remained stable once VPDs exceeded the BP (Fig. 5 I,L). Across the entire  
380 range of VPDs, iWUE did not significantly differ between genotypes. However, *Az34*  
381 had a significantly ( $P < 0.001$ ) lower iWUE after the BP (1.9 kPa). Thus, the importance  
382 of ABA in determining iWUE of these genotypes varied according to the VPD.

383

384 **Differences in whole plant gas exchange in response to ABA in an ABA-**  
385 **deficient mutant**

386 Before applying ABA and while at 2.5 kPa VPD,  $T_{IRGA}$  and  $A_n$  were 13% higher  
387 in WT than *Az34* plants, while  $iWUE$  did not significantly differ between genotypes  
388 (Table 2). Foliar ABA application reduced  $T_{IRGA}$  within 5 min in both the ABA-deficient  
389 mutant *Az34* and WT barley, with  $T_{IRGA}$  stabilising after 1 h (Supplemental Fig. 7).  
390 Whole plant  $T_{IRGA}$  decreased by 40% and 23% in *Az34* and WT plants respectively  
391 (Table 2); with the response almost significantly greater in *Az34* ( $P = 0.053$  for  
392 genotype x ABA interaction). Interestingly, ABA application did not significantly affect  
393  $A_n$  of WT plants, but decreased  $A_n$  by 30% in *Az34*. ABA treatment increased  $iWUE$   
394 similarly in both genotypes (no significant genotype x ABA interaction). Taken  
395 together, whole plant gas exchange of the ABA-deficient mutant *Az34* was more  
396 responsive than WT plants to foliar ABA application.

397

398 **Leaf-level measurements**

399 To further investigate the mechanisms by which ABA limits photosynthesis, the  
400 flag leaves of *Az34* and WT plants were sprayed with ABA (Table 3). Stomatal  
401 conductance and leaf internal  $CO_2$  concentration ( $C_i$ ) were 50% and 15% higher,  
402 respectively, in *Az34* than WT plants in the greenhouse prior to applying ABA.  
403 Following ABA application, both  $g_s$  and  $C_i$  decreased, more severely in *Az34* plants  
404 as indicated by significant ( $P < 0.001$  and  $P < 0.009$ ) genotype x ABA interactions  
405 (Table 3). *Az34* had ca. 50% less total soluble protein (TSP) and Rubisco  $V_t$  than WT  
406 plants. *Az34* showed higher Rubisco activation state than WT prior to ABA application,  
407 with the opposite observed after ABA application as indicated by the significant ( $P <$

408 0.001) genotype x ABA interaction (Table 3). ABA had no significant effect ( $P > 0.05$ )  
409 on TSP or Rubisco  $V_t$  for either genotype. However, while activation states were not  
410 affected by the ABA treatment in WT plants, *Az34* significantly reduced the activation  
411 state by ca. 25% (Table 3). Flag leaf ABA concentration of *Az34* was approximately  
412 half of the value in WT plants before spraying, and ABA application increased leaf  
413 [ABA] of both genotypes by 6-7 fold (Supplemental Table 5). Taken together, stomatal  
414 and photosynthetic responses of *Az34* were more responsive to exogenous ABA  
415 spraying, despite similar proportional changes in foliar ABA accumulation.

416

## 417 **DISCUSSION**

### 418 **The whole plant chamber can identify genetic diversity in gas exchange**

419 A whole plant gas exchange chamber was adapted to study plant  $T_{IRGA}$ ,  $A_n$  and  $iWUE$   
420 responses to changing VPD. The findings with wheat and barley genotypes support  
421 the idea that the chamber enables a robust assessment of these responses. A  
422 previous study demonstrated that some wheat genotypes restrict  $T_r$  at high VPD, such  
423 as cv. Drysdale [16], here we also show that this response significantly improves  $iWUE$   
424 since photosynthesis is not limited above the BP. This reinforces the idea that  $iWUE$   
425 can be improved by including the restricted transpiration trait at high VPD in those  
426 wheat genotypes that do not show it because  $A_n$  is not limited by VPD, making this an  
427 effective strategy to implement in breeding programs for drought-prone environments  
428 in elite plants [49-50].

429 The whole plant gas exchange system was developed for phenotyping whole  
430 plant  $iWUE$  at different VPDs, and identified genetic differences. At a single VPD,  
431 genotypic differences in  $A_n$  correlated with single-leaf measurements done in field

432 conditions in a previous experiment [36]. Nevertheless, at that specific VPD, a similar  
433  $T_{IRGA}$  was found between Drysdale and Krichauff, in contrast with the results when  
434 comparing such genotypes under different VPD, reinforcing the importance of the VPD  
435 response curves in ranking  $T_{IRGA}$ . It is important to note that plants were exposed to  
436 high VPD by maintaining air temperatures lower than 30°C, which does not limit wheat  
437 photosynthesis [51,52]. However, under natural conditions, high VPD and  
438 temperatures occur together, with inhibition of  $A_n$  by high VPD attributed to  
439 excessively high temperatures [53]. Moreover, under non-steady state conditions, high  
440 VPD can constrain photosynthetic induction: the time required to reach the maximum  
441  $A_n$  after the transition from low to high light [54]. Taken together, our results show that  
442 restricting  $T_{IRGA}$  at high VPDs at an optimal temperature range and under steady-state  
443 conditions does not affect carbon assimilation in commercial wheat and barley  
444 cultivars. It is essential to understand the physiological mechanisms regulating these  
445 responses.

446

#### 447 **Determining the role of ABA in VPD responses**

448 Previous measurements at whole plant level using gravimetric methods [8, 41]  
449 have implicated ABA in regulating cereal transpiration under varying evaporative  
450 demands. Similarly, transpiration of the ABA-deficient barley mutant *Az34* was  
451 unrestricted at high VPDs, but unexpectedly,  $A_n$  was limited (Fig. 4K; Supplemental  
452 Table 3). Single-leaf measurements were required to confirm the mechanistic  
453 response to the reduction of photosynthesis in ABA-deficient plants. Despite higher  
454 intercellular  $CO_2$  concentrations due to greater stomatal opening, *Az34* had a lower  
455 Rubisco activity (ca. 70% reduction compared to WT plants). Since *Az34* is nitrate-

456 reductase deficient [40], plants are expected to be N limited with approximately half  
457 the total soluble protein content compared to WT plants (Table 3). Thus, the limited  
458 biomass of *Az34* not only results from its inability to control water loss under moderate-  
459 high VPD [41, 55], which induces leaf water deficit, but also from reduced Rubisco  
460 carboxylation that lowers photosynthesis.

461 To further demonstrate that dynamic whole plant responses can be detected  
462 with our system, ABA was sprayed on the leaves [12, 38]. Exogenous ABA application  
463 decreased Tr by ca. 25% in WT plants but even more so in *Az34* (by 40%), indicating  
464 greater stomatal sensitivity of the ABA-deficient mutant. These differences in whole  
465 plant transpiration sensitivity to ABA were confirmed in flag leaves (Table 3). Several  
466 ABA-deficient mutants in *Arabidopsis* (*aba2-11*, *nced3 nced 5*, *aba1-1*, *aba4-3*, *aao3-*  
467 *2*, *aba3-1*) and other species (*wilty* pea and *flacca* tomato) were described as  
468 hypersensitive to exogenous ABA application [35], attributed to a higher pre-treatment  
469 gs. Further work is required to investigate possible feedback regulation of genes for  
470 ABA sensitivity by ABA status in ABA-deficient mutants.

471 The mechanisms by which exogenous ABA limits photosynthesis remain under  
472 debate. While stomatal closure after ABA application decreases  $C_i$  even in ABA-  
473 deficient mutants ([38]; Table 3 here), the ABA molecule has been suggested to bind  
474 to the Rubisco active site blocking Rubisco activity [57]. While foliar ABA spraying did  
475 not affect photosynthesis of WT plants (Table 2),  $A_n$  was decreased by 30% in *Az34*,  
476 as in a previous comparison of WT and ABA-deficient tomatoes (*flacca* mutant) grown  
477 under non-saturating light (Bradford et al. 1983). ABA application decreased Rubisco  
478 activation state of *Az34* flag leaves but not WT leaves (Table 2). Similar to *in vitro* ABA  
479 experiments [57], activation of such plants might be disrupted by the ABA molecule.  
480 Alternatively, *Az34* may have higher  $CO_2$  availability under standard conditions in the

481 greenhouse (before ABA application). The larger decreases in  $g_s$  and  $C_i$  observed in  
482 *Az34* after ABA application may deactivate Rubisco because of the limited  $CO_2$   
483 availability, thereby decreasing photosynthesis. Whether such limitations occur  
484 because stomatal and mesophyll conductance are co-ordinated [58], or due to a  
485 mechanistic constraint of ABA on Rubisco activity, is still unknown. In either case, the  
486 lower Rubisco activity of *Az34* makes its photosynthesis more vulnerable to  
487 environmental constraints, such as high VPD, than WT plants.

488

## 489 **CONCLUSIONS**

490 Our chamber was designed, built and operated to evaluate whole plant  $A_n$ ,  
491  $T_{IRGA}$  and  $iWUE$  under increasing evaporative demand in small-grain cereals. This  
492 instrumentation is sufficiently precise to detect genetic differences in plant responses.  
493 In wild-type genotypes, photosynthesis was not restricted by VPD “*per se*”, even  
494 though some genotypes restricted  $T_{IRGA}$  under high VPD, which is of direct interest to  
495 plant breeders seeking to increase  $iWUE$ . Furthermore, ABA-deficient barley  
496 responded more sensitively to exogenous ABA application, with greater transpirational  
497 restriction and decreased Rubisco activation state. Photosynthesis of ABA-deficient  
498 barley plants was also limited at high VPD, likely due to reduced Rubisco activity.

499

## 500 **DECLARATIONS**

### 501 **Author’s contribution**

502 ICD designed the research with input from all authors; IJ developed the system and  
503 conducted the experiments with initial input from SAR; IJ analyzed the results; all

504 authors contributed to interpret the results; IJ and ICD wrote the manuscript with  
505 contributions from SAR, SHT, MAJP, ECS. All authors read and approved the final  
506 manuscript.

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511 the physiological interpretation of the results obtained.

### 512 **Competing interests**

513 The authors declare that they have no competing interests.

### 514 **Availability of data and materials**

515 All data generated or analyzed during this study are included in this published article.  
516 The datasets used and analyzed during the current study are available from the  
517 corresponding author on reasonable request.

### 518 **Consent for publication**

519 Not applicable.

### 520 **Ethics approval and consent to participate**

521 Not applicable.

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526

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