

1 **Short title:** Systemic ROS signaling during stress

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4 **Vascular and non-vascular transmission of systemic reactive**  
5 **oxygen signals during wounding and heat stress**

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14 **One-sentence summary:** In addition to vascular bundles, mesophyll cells can mediate the ROS  
15 wave during systemic responses to wounding or heat stress in Arabidopsis.

16 **Author Contributions:** S.I.Z. performed the experiments. S.I.Z. and R.M. designed the  
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24

25 **ABSTRACT**

26 Sensing of heat, high light (HL), or mechanical injury by a single leaf of a plant results in the  
27 activation of different systemic signals that reach systemic tissues within minutes and trigger  
28 systemic acquired acclimation (SAA) or systemic wound responses (SWRs), resulting in a  
29 heightened state of stress readiness of the entire plant. Among the different signals associated  
30 with rapid systemic responses to stress in plants are electric, calcium and reactive oxygen species  
31 (ROS) waves. These signals propagate from the stressed or injured leaf to the rest of the plant  
32 through the plant vascular bundles, and trigger SWRs and SAA in systemic tissues. However,  
33 whether they can propagate through other cell types, and whether or not they are interlinked,  
34 remain open questions. Here we report that in response to wounding or heat stress (HS), but not  
35 HL stress, the ROS wave can propagate through mesophyll cells of *Arabidopsis thaliana*.  
36 Moreover, we show that propagation of the ROS wave through mesophyll cells during these  
37 stresses is sufficient to restore SWR and SAA transcript accumulation in systemic leaves, as well  
38 as SAA to HS (but not HL). We further show that propagation of the ROS wave through  
39 mesophyll cells could contribute to systemic signal integration during HL&HS stress  
40 combination. Our findings reveal that the ROS wave can propagate through tissues other than the  
41 vascular bundles of plants, and that different stresses can trigger different types of systemic  
42 signals that propagate through different cell layers and induce stress-specific systemic responses.

43

44 **Key words:** Abiotic stress, *Arabidopsis thaliana*, Heat stress, High light stress, Mesophyll, ROS  
45 wave, Systemic signaling, Vascular bundles, Wounding.

46

47 **Abbreviations:** HL, high light; HS, heat stress; RBOHD, respiratory burst oxidase homolog D;  
48 ROS, reactive oxygen species; SAA, Systemic acquired acclimation; SWR, systemic wound  
49 response.

50

## 51 INTRODUCTION

52 In response to different abiotic stresses plants mount an acclimation response that counters the  
53 adverse effects of stress on plant metabolism, reproduction, and overall survival (Zhu, 2016;  
54 Kollist et al., 2019; Cheung et al., 2020). This response is triggered upon perception of stress at  
55 the tissues immediately subjected to stress (termed local tissues), as well as in other tissues of the  
56 plant that have not yet experienced the stress (termed systemic tissues). The perception of stress  
57 at the local tissues activates therefore a signal transduction process that links the different tissues  
58 (local to all systemic tissues) over long distances, sometime spanning the entire length of the  
59 plant (*e.g.*, Miller et al., 2009; Szechyńska-Hebda et al., 2010; Christmann et al., 2013; Choi et  
60 al., 2014; Gilroy et al., 2016; Guo et al., 2016; Choi et al., 2017; Choudhury et al., 2018;  
61 Devireddy et al., 2018; Takahashi et al., 2018; Toyota et al., 2018; Fichman et al., 2019; Wang et  
62 al., 2019; Zandalinas et al., 2019; Devireddy et al., 2020b; Devireddy et al., 2020a; Farmer et al.,  
63 2020; Fichman and Mittler, 2020). This process is termed systemic signaling, and the  
64 acclimation of systemic tissues to stress, upon perception of the systemic signal, is called  
65 systemic acquired acclimation (SAA; Karpinski et al., 1999). A similar systemic signaling  
66 process occurs in plants upon wounding of a local leaf, and this process is termed systemic  
67 wound response (SWR; Walker-Simmons et al., 1984). During SAA or SWR, many different  
68 abiotic stress- or wound-response transcripts and hormones that rapidly accumulate in the local  
69 leaf upon stress or wounding also accumulate within minutes in the systemic tissues, and these  
70 transcripts and hormones are thought to mediate SAA or SWR at the systemic tissues (*e.g.*,  
71 Galvez-Valdivieso et al., 2009; Miller et al., 2009; Suzuki et al., 2013; Zandalinas et al., 2019;  
72 Fichman et al., 2020b). Although the process of SAA or SWR can be easily traced back to some  
73 of the regulatory transcripts and hormones that accumulate in systemic tissues during stress, how  
74 the systemic signal initiating at the local leaf and reaching the systemic tissues is propagated, and  
75 what is its nature, are still ongoing subjects of active research (*e.g.*, Fichman et al., 2020a;  
76 Fichman and Mittler, 2020). Among the main candidates for systemic signals mediating SAA or  
77 SWRs are electric, calcium, reactive oxygen species (ROS), and hydraulic pressure waves  
78 (Miller et al., 2009; Christmann et al., 2013; Mousavi et al., 2013; Choi et al., 2014; Nguyen et  
79 al., 2018; Toyota et al., 2018; Shao et al., 2020).

80 Because plants lack a true nervous system that connects different tissues, systemic signals that  
81 travel from the local tissue, initially subjected to stress, to the entire plant are transmitted by cell-  
82 to-cell signaling events that involve changes in calcium, membrane potential and ROS (Fichman  
83 and Mittler, 2020). It is thought that during this process the different cells along the path of the  
84 cell-to-cell signaling chain are being activated one-by-one (similar to a domino effect) starting at  
85 the initial (local) tissue and ending at the systemic tissue, and that this activation process  
86 propagates and maintains the different systemic signals. This concept was initially proposed as a  
87 way to transmit ROS signals over long distances in plants (Miller et al., 2009; Mittler et al.,  
88 2011), and was later adopted for explaining calcium and other systemic signals (Choi et al.,  
89 2014, 2017). According to this model, each cell along the cell-to-cell path that transmits the  
90 signal starts to actively generate ROS upon sensing that the cell preceding it in the chain is  
91 producing ROS. It was found that in *Arabidopsis thaliana* the ROS produced by each cell during  
92 this process is generated by the respiratory burst oxidase homolog D (RBOHD) protein and that  
93 this process is controlled by calcium-dependent activation of RBOHD (Fichman and Mittler,  
94 2020; Fichman et al., 2021). The ROS being used as a systemic signal, most likely H<sub>2</sub>O<sub>2</sub> (Miller  
95 et al., 2009), is therefore actively generated by each cell along the path of the signal, as opposed  
96 to being made in the local tissue and somehow transported over long distances (Mittler et al.,  
97 2011).

98 Recently, wound-induced systemic cell-to-cell electric and calcium signals were shown to be  
99 dependent on the function of glutamate receptor-like (GLR) calcium channels expressed at the  
100 vascular bundles of *Arabidopsis*, and a double mutant for *glr3.3;glr3.6* was shown to be deficient  
101 in wound-induced systemic signaling (Mousavi et al., 2013; Nguyen et al., 2018; Toyota et al.,  
102 2018; Shao et al., 2020). In contrast, systemic cell-to-cell signaling, and SAA to high light (HL)  
103 or heat stress (HS) were found to be dependent on ROS produced in each cell along the path of  
104 the signal by RBOHD and/or RBOHF (Miller et al., 2009; Fichman et al., 2019; Zandalinas et  
105 al., 2020b). At least in response to HL stress, this process was also found to occur at the vascular  
106 bundles of *Arabidopsis* (Zandalinas et al., 2020b). A new study has now revealed that GLR3.3  
107 and/or GLR3.6 are not absolutely required for HL-induced systemic ROS signaling, and that the  
108 systemic signal mediating SAA to HL stress in *Arabidopsis* requires a coordinated function of  
109 plasmodesmata (PD) proteins (*i.e.*, plasmodesmata-localized proteins 1 and 5; PDLP1 and  
110 PDLP5) and RBOHD (Fichman et al., 2021). It was further found that RBOHD-produced ROS

111 opens PD pores between cells and facilitates cell-to-cell transport during this process, suggesting  
112 that enhancing transport through PDs is one possible role for ROS during systemic cell-to-cell  
113 signaling in plants (Fichman et al., 2021). In addition, aquaporins such as PIP2;1 and calcium-  
114 permeable channels, such as cyclic nucleotide-gated calcium channel 2 (CNGC2), and  
115 mechanosensitive small conductance-like (MSL) channels 2 and 3 were found to be involved in  
116 this process (Fichman et al., 2021). Moreover, in response to wounding the systemic ROS signal  
117 was shown to induce a systemic redox signal (wave) that propagated throughout the plant within  
118 minutes (Fichman and Mittler, 2021).

119 A recent study has also revealed that in contrast to the local application of HL or HS to a single  
120 leaf of Arabidopsis, or the co-application of HL and HS to the same leaf (HL+HS), the co-  
121 application of HL and HS to two different leaves of the same plant (HL&HS) resulted in a  
122 stronger ROS wave response (Zandalinas et al., 2020a). It was further found that the plant  
123 hormone jasmonic acid (JA) suppresses the activation of the ROS wave in local leaves  
124 simultaneously subjected to a combination of HL and HS (HL+HS; Zandalinas et al., 2020a).  
125 Although the ROS wave was found to propagate through the vascular bundles of Arabidopsis  
126 during systemic responses to HL stress (Zandalinas et al., 2020b), it is unknown at present  
127 whether it propagates through the same plant tissues during other stresses, such as HS or  
128 wounding. Finding, for example, that the ROS wave propagates through other plant tissues  
129 during HS, could provide a potential explanation to the stronger ROS wave signal observed  
130 under conditions of HL&HS (Zandalinas et al., 2020a). In addition, it could provide initial  
131 evidence for the propagation of rapid systemic signals outside the vascular bundles of plants.

132 To identify the plant tissues that mediate RBOHD-dependent systemic ROS signal propagation  
133 during responses to HS or wounding, we used the *rbohD* transgenic lines we previously  
134 developed to study the propagation of the ROS wave during HL stress (Zandalinas et al., 2020b).  
135 Our findings reveal that in contrast to RBOHD-dependent systemic responses to HL stress, that  
136 were exclusively mediated through the vascular bundles of Arabidopsis (Zandalinas et al.,  
137 2020b), RBOHD-dependent systemic signaling during HS (Zandalinas et al., 2020a), or  
138 wounding (Miller et al., 2009; Fichman et al., 2019; Fichman and Mittler, 2021), are mediated  
139 through the vascular bundles and/or mesophyll cells. We further show that propagation of the  
140 ROS wave through mesophyll cells could contribute to the stronger systemic ROS signal

141 observed in plants subjected to HL and HS simultaneously applied to two different leaves  
142 (HL&HS; Zandalinas et al., 2020a). Our findings provide direct evidence for the propagation of  
143 rapid systemic ROS signals through tissues other than the vascular bundles of Arabidopsis.

## 144 **RESULTS**

### 145 **Vascular bundles or mesophyll cells can mediate the ROS wave during the systemic** 146 **response of Arabidopsis to wounding**

147 To identify the plant tissues that transmit RBOHD-dependent systemic **signals** (*i.e.*, the ROS  
148 wave; Miller et al., 2009) in response to a local application of wounding, we used the different  
149 transgenic lines we previously developed of *rbohD*, in which *RBOHD* was expressed under its  
150 native promoter or different tissue-specific promoters (Zandalinas et al., 2020b). These lines  
151 were previously characterized for their ROS wave propagation, SAA and *Zat12* expression in  
152 response to a local application of HL stress, and the localization and stable expression of the  
153 RBOHD protein in their different tissues was confirmed using GFP-RBOHD fusions **driven by**  
154 **the tissue-specific promoters** (Zandalinas et al., 2020b). In our analysis we included wild-type,  
155 *rbohD* null mutants, and *rbohD* mutants in which *RBOHD* was expressed under its native  
156 promoter, or epidermis-, mesophyll- xylem parenchyma-, phloem- or bundles sheath-specific  
157 promoters (Zandalinas et al., 2020b; **Supplementary Figure S1**). All plants were wounded on a  
158 single local leaf and local and systemic ROS levels were imaged in whole-plants grown in soil  
159 using the new live-imaging method we developed to image ROS (Fichman et al., 2019;  
160 Zandalinas et al., 2020a; Zandalinas et al., 2020b). As shown in Figure 1 **and Supplementary**  
161 **Figure S2**, wound-induced systemic ROS accumulation was suppressed in the *rbohD* mutant and  
162 this suppression was complemented to wild type levels by expression of *RBOHD* in the *rbohD*  
163 mutant using its native promoter. Expressing the RBOHD protein in *rbohD* plants using the  
164 mesophyll-, xylem parenchyma- or phloem-specific promoters also complemented the systemic  
165 accumulation of ROS to wild type levels in the *rbohD* mutant in response to wounding. In  
166 contrast, as shown in Supplementary **Figures S2, S3**, as well as previously reported (Zandalinas  
167 et al., 2020b), in response to a local application of HL stress, expression of the RBOHD protein  
168 in *rbohD* plants using the native promoter of *RBOHD*, or using the xylem parenchyma- or  
169 phloem-specific promoters (but not the mesophyll-specific promoter), complemented the  
170 systemic accumulation of ROS in *rbohD* mutants to wild type levels in response to HL. These

171 finding reveal that in response to a local wounding treatment, the ROS wave can propagate  
172 through the vascular bundles, or mesophyll cells of Arabidopsis.

173

### 174 **Vascular bundles or mesophyll cells can mediate the ROS wave during the systemic** 175 **response of Arabidopsis to heat stress**

176 As shown in Figure 2 and Supplementary Figure S2, a similar result to that shown in Figure 1  
177 was obtained when a local Arabidopsis leaf was subjected to HS. Thus, similar to the local  
178 application of wounding (Figure 1), but different from the local application of HL  
179 (Supplementary Figures S2, S3; Zandalinas et al., 2020b), expression of the RBOHD protein in  
180 *rbohD* plants using its native promoter, or using the mesophyll-, xylem parenchyma- or phloem-  
181 specific promoters, complemented the systemic accumulation of ROS in *rbohD* mutants to wild  
182 type levels in response to a local application of HS. The findings shown in Figures 1, 2, and  
183 Supplementary Figure S2, reveal therefore that unlike rapid systemic ROS responses to HL, that  
184 could only be complemented to wild type levels in the *rbohD* mutant by expressing the RBOHD  
185 protein in xylem parenchyma or phloem cells (Supplementary Figures S2, S3; Zandalinas et al.,  
186 2020b), tissues limited in their localization to the vascular bundles, systemic ROS signals (*i.e.*,  
187 the ROS wave) to wounding or HS can be mediated by RBOHD protein found in mesophyll  
188 cells, that are primarily localized outside the vascular bundles of plants.

189

### 190 **Complementing the ROS wave by expression of RBOHD in mesophyll cells restores SAA-** 191 **and SWR-associated transcript expression in systemic leaves in response to a local HS or** 192 **wounding treatment**

193 Complementing the ROS wave by expression of *RBOHD* in mesophyll cells (Figures 1, 2, and  
194 Supplementary Figures S1, S2) might or might not complement the expression of systemic  
195 transcripts previously associated with SAA or SWR in response to a local application of HS or  
196 wounding, respectively. Complementation of *RBOHD* expression in the *rbohD* mutant using the  
197 xylem parenchyma- or phloem- (but not mesophyll-) specific promoters restored the expression  
198 of the *Zat12* SAA and SWR gene in response to local application of HL stress (measured using  
199 *Zat12:luciferase;rbohD* double mutants complemented with the different tissue-specific *RBOHD*

200 transformation vectors; Zandalinas et al., 2020b). Because *Zat12* reporter plants might not be a  
201 good experimental tool to study stress-specific responses to HS, HL or wounding (*Zat12* is  
202 expressed in response to HL or wounding; Miller et al., 2009), we elected to study the expression  
203 of different wounding-, HS-, or HL-specific transcripts in the different lines shown in Figures 1  
204 and 2 in response to a local application of HS, wounding, or HL using quantitative RT-PCR  
205 (qPCR). We chose the transcripts and timing for this analysis based on our previous RNA-Seq  
206 studies of systemic signaling in response to HL and/or HS (Suzuki et al., 2013; Zandalinas et al.,  
207 2019; Fichman et al., 2020b; Zandalinas et al., 2020a), as well as based on studies of systemic  
208 wound responses using transcriptomics and qPCR analyses (Suzuki et al., 2013; Toyota et al.,  
209 2018). As shown in Figure 3A, expression of the wound-response transcripts *JAZ5* and *JAZ7* was  
210 enhanced in local and systemic leaves of wild type plants upon local wounding. In contrast, in  
211 response to the same treatment, the expression of these transcripts was suppressed in systemic  
212 (but not local) leaves of the *rbohD* mutant. Complementation of *RBOHD* expression with the  
213 *RBOHD* native promoter, or the mesophyll-, xylem parenchyma-, or phloem-specific promoters  
214 restored the systemic expression of *JAZ5* and *JAZ7* in response to a local wounding treatment. In  
215 contrast, complementation of *RBOHD* expression with the bundle sheath- or epidermis-specific  
216 promoters failed to restore the systemic expression of *JAZ5* and *JAZ7* to wild type levels in  
217 response to the local wounding treatment. These findings reveal that complementing the ROS  
218 wave by expression of *RBOHD* in mesophyll, xylem parenchyma or phloem cells of the *rbohD*  
219 mutant was sufficient to restore some SWR-specific transcript expression in response to a local  
220 wounding treatment.

221 To test the effect of restoring *RBOHD* expression in the different tissues on SAA responses to  
222 HS, we studied the expression of *Rap2.4* and *ERF2*, two transcripts previously associated with  
223 SAA to HS (Suzuki et al., 2013; Zandalinas et al., 2020a), in local and systemic leaves of the  
224 different wild type, *rbohD* and *rbohD*-complemented lines, in response to a local HS treatment.  
225 As shown in Figure 3B, expression of the HS-response transcripts *Rap2.4* and *ERF2* was  
226 enhanced in local and systemic leaves of wild type plants upon a local HS treatment. In contrast,  
227 in response to the same treatment, the enhanced expression of these transcripts was blocked in  
228 systemic and suppressed in local leaves of the *rbohD* mutant. Complementation of *RBOHD*  
229 expression with the *RBOHD* native promoter, or the mesophyll-, xylem parenchyma-, or phloem-  
230 specific promoters restored the systemic expression of *Rap2.4* and *ERF2* in response to a local



231 HS treatment. In contrast, complementation of *RBOHD* expression with the bundle sheath- or  
232 epidermis-specific promoters did not restore the systemic expression of *Rap2.4* and *ERF2* to  
233 wild type levels in response to a local HS treatment. These findings reveal that similar to the  
234 response of *JAZ5* and *JAZ7* to wounding (Figure 3A), restoring the ROS wave by expression of  
235 *RBOHD* in mesophyll, xylem parenchyma or phloem cells was sufficient to restore some SAA  
236 transcript expression in systemic leaves in response to a local HS treatment.

237 To study whether a similar effect would occur in complemented *rbohD* plants subjected to a  
238 local treatment of HL, we studied the expression of *MYB30* and *ZHD5*, two transcripts associated  
239 with the SAA response of Arabidopsis to HL stress (Zandalinas et al., 2019; Fichman et al.,  
240 2020b; Zandalinas et al., 2020a). As shown in Figure 3C, similar to *Zat12* expression in the  
241 different *rbohD*-complemented lines (Zandalinas et al., 2020b), complementation of *RBOHD*  
242 expression in xylem parenchyma or phloem (but not mesophyll) cells of the *rbohD* mutant  
243 supported the systemic expression of *MYB30* and *ZHD5* in response to a local treatment of HL  
244 stress. Complementation of *RBOHD* expression in mesophyll cells of the *rbohD* mutant did  
245 however result in enhanced local (but not systemic) expression of *MYB30* and *ZHD5* (Figure  
246 3C), demonstrating that local leaves of these plants were able to sense the HL stress but were  
247 unable to initiate the systemic ROS signal in response to it. Taken together, the results presented  
248 in Figures 1-3 and Supplementary Figures S1-S3 reveal that complementing the expression of  
249 *RBOHD* in the mesophyll, xylem parenchyma or phloem cells of the *rbohD* mutant restores not  
250 only the ROS wave, but also the expression of certain systemic transcripts specific to wounding  
251 or HS. In contrast, complementing the expression of *RBOHD* in mesophyll cells of the *rbohD*  
252 mutant did not complement the ROS wave or systemic HL-specific SAA transcripts in response  
253 to a local application of HL stress (Figure 3 and Supplementary Figure S3).

254

### 255 **Complementing the ROS wave by expression of *RBOHD* in mesophyll cells restores local** 256 **HS-induced SAA**

257 Complementing the expression of *RBOHD* in the xylem parenchyma or phloem cells of the  
258 *rbohD* mutant restored SAA to HL (Supplementary Figure S4; Zandalinas et al., 2020b).  
259 Although we do not have a biological assay for SAA during SWR, aside from measuring  
260 systemic wound-induced transcript expression as shown in Figure 3, an assay for SAA to HS was

261 previously reported (Suzuki et al., 2013; Zandalinas et al., 2020a). We therefore used this assay  
262 to study whether restoring *RBOHD* expression in mesophyll cells could restore SAA to HS of the  
263 *rbohD* mutant. As shown in Figure 4, complementing the expression of *RBOHD* in the *rbohD*  
264 mutant using its native promoter, or the mesophyll-, xylem parenchyma-, or phloem-specific  
265 promoters restored SAA to HS. In contrast, complementing the expression of *RBOHD* in the  
266 *rbohD* mutant using the mesophyll-specific promoter failed to restore SAA to HL  
267 (Supplementary Figure S4; Zandalinas et al., 2020b). The findings presented in Figures 1-4 and  
268 Supplementary Figures S1-S4 reveal therefore that expression of *RBOHD* in mesophyll cells can  
269 restore the ROS wave, systemic transcript expression, and SAA to HS (but not HL stress) in the  
270 *rbohD* mutant.

271

#### 272 **Could expression of *RBOHD* in mesophyll cells contribute to the stronger systemic ROS** 273 **signal observed in plants subjected to HL&HS?**

274 We previously reported that HS and HL, when applied to two different leaves of the same  
275 Arabidopsis plant (HL&HS), result in a stronger ROS wave response compared to HS or HL  
276 applied to a single leaf, or to the same leaf (HL+HS; Zandalinas et al., 2020a). Our current  
277 findings that in response to HS the ROS wave could be mediated through mesophyll, xylem  
278 parenchyma, and/or phloem cells (Figures 2-4), but in response to HL it could only be mediated  
279 through xylem parenchyma and/or phloem cells (Supplementary Figures S2-S4; Zandalinas et  
280 al., 2020b), might provide a potential explanation to this phenomena. In response to HL and HS  
281 applied to two different leaves (HL&HS), the systemic ROS wave might be stronger because it  
282 would propagate through an additional cell layer (mesophyll, contributed by the HS treatment).  
283 This could not occur of course when the two stresses are applied to the same leaf because under  
284 these conditions the ROS wave induced by HL+HS applied to the same leaf is suppressed by JA  
285 (Zandalinas et al., 2020a). To test whether the ROS wave could propagate through mesophyll  
286 cell layers during HL&HS combination, we compared the intensity of the ROS wave between  
287 wild type, *rbohD*, and *rbohD* in which *RBOHD* expression was complemented at the mesophyll  
288 or phloem cells, subjected to a HL&HS treatment (Figure 5 and Supplementary Figure S5). As  
289 shown in Figure 5, compared to wild type plants, the ROS wave was suppressed in *rbohD* plants  
290 subjected to the HL&HS treatment. Complementation of *RBOHD* expression with *RBOHD*

291 expressed under the control of its native promoter, or a phloem specific promoter (that could  
292 restore HS- or HL-response ROS wave functions from the two different leaves; Figures 2-4 and  
293 Supplementary Figures S2, S3 and S5; Zandalinas et al., 2020b) restored the ROS wave to its  
294 high level of expression. In contrast, complementation of the *rbohD* mutant with *RBOHD*  
295 expressed under the mesophyll- specific promoter (that could only restore HS-, but not HL-  
296 response ROS wave functions from the HL-treated leaf; Figures 2-4 and Supplementary Figures  
297 S2, S3 and S5; Zandalinas et al., 2020b), could not restore the ROS wave to its maximal  
298 intensity. These finding demonstrate that under conditions of HL&HS at least part of the ROS  
299 wave that spreads throughout the plant (originating from the HS-treated leaf) could be mediated  
300 through mesophyll cells. Complementation of *RBOHD* expression with *RBOHD* expressed under  
301 the control of the phloem-specific promoter was nonetheless sufficient to restore the ROS wave  
302 to wild type or *rbohD* mutant complemented with *RBOHD* under its native promoter levels  
303 (Figure 5), suggesting that in wild type plants transmission of the ROS wave signal through  
304 phloem cells is sufficient to cause a higher signal during HL&HS combination.

305

## 306 DISCUSSION

307 Abiotic, mechanical injury, and biotic stresses trigger a rapid systemic signal transduction  
308 process that activates different acclimation and defense mechanisms in systemic tissues within  
309 minutes of stress sensing at the local tissues (Fichman et al., 2019; Kollist et al., 2019; Fichman  
310 and Mittler, 2020). Up until now, the systemic electric, calcium and ROS waves, triggered by  
311 wounding or HL stress, were shown to be mediated through the vascular bundles of plants  
312 (Mousavi et al., 2013; Nguyen et al., 2018; Toyota et al., 2018; Farmer et al., 2020; Shao et al.,  
313 2020; Zandalinas et al., 2020b). Here, we present evidence that in addition to vascular bundles,  
314 mesophyll cells can also mediate the systemic ROS wave in response to a local treatment of  
315 wounding or HS (Figures 1-6). Mesophyll cells are not typically considered part of the vascular  
316 bundles of plants and are found within leaves and stems as cell layers that connect the vascular  
317 tissues to the epidermis, stomata and/or other leaf/stem structures and cell types. Because the  
318 ROS wave propagates from cell-to-cell via mechanisms that require apoplastic and symplastic  
319 connectivity between cells (Miller et al., 2009; Fichman et al., 2021), and mesophyll cells are  
320 connected with each other via PD and/or their shared apoplastic microenvironment, as well as

321 express *RBOHD* under controlled growth conditions (Supplementary Figure S1; Zandalinas et  
322 al., 2020b), the basic mechanisms that allow the ROS wave to propagate from cell-to-cell  
323 through mesophyll cell layers appear to be present. In contrast, GLR3.3 and/or GLR3.6 that are  
324 required for rapid wound-response systemic signaling are not thought to be localized to  
325 mesophyll cells (they are thought to be exclusively localized to the xylem parenchyma and  
326 phloem cells; Mousavi et al., 2013; Nguyen et al., 2018; Toyota et al., 2018; Shao et al., 2020).  
327 A recent study has shown that GLR3.3 and/or GLR3.6 are not absolutely required for the ROS  
328 wave to propagate in response to a local treatment of HL stress (Fichman et al., 2021). Taking  
329 this study into consideration, it is plausible that the ROS wave will propagate through tissues that  
330 do not express GLR3.3 and/or GLR3.6, possibly using other calcium-permeable channels such as  
331 CNGCs or MSLs (Fichman et al., 2021). Having many of the required proteins and physical  
332 connections/proximity required for ROS cell-to-cell signals to function, support the possibility  
333 that the ROS wave can propagate through layers of mesophyll cells that are outside the vascular  
334 bundles (Figure 6).

335 Considering the extensive literature and established role of chloroplasts in the perception of light  
336 stress, as well as ROS production (Karpinski et al., 1999; Mittler, 2002), it is somewhat  
337 surprising that perception of light stress in *rbohD* mutants expressing RBOHD in mesophyll cells  
338 does not trigger the systemic ROS wave (Supplementary Figure S3; Zandalinas et al., 2020b).  
339 Although local leaves of *rbohD/pCAB3::GFP-RbohD* plants express MYB30 and ZHD5 in  
340 response to HL stress, showing that they can perceive the stress, they are nevertheless unable to  
341 trigger the systemic ROS signal and cause accumulation of these transcripts in systemic leaves  
342 (Figure 3 and Supplementary Figure S3). One possible explanation to this finding stem from  
343 recent studies showing that HL-induced ROS in Arabidopsis leaves and bundle sheath cells of  
344 rice requires RBOH proteins (Devireddy et al., 2020b; Xiong et al., 2021). It is therefore possible  
345 that triggering of the systemic ROS signal during light stress requires RBOH present in vascular  
346 cells and that this process is independent of ROS accumulation in chloroplasts of mesophyll  
347 cells. Further studies are required to address this intriguing possibility.

348 In addition to showing that the ROS wave can propagate outside the vascular bundles of  
349 Arabidopsis (Figures 1 and 2 and Supplementary Figure S2), supporting systemic wound- and  
350 HS-induced transcript expression in systemic leaves (Figure 3) and mediating SAA to HS

351 (Figure 4), our findings further highlight the interesting possibility that different stresses, *e.g.*,  
352 HS, HL and wounding, trigger different types of systemic waves that propagate through different  
353 tissues, and could even be spatially separated from each other. For example, complementing  
354 *RBOHD* expression in mesophyll cells of the *rbohD* mutant can complement systemic responses  
355 to wounding (Figure 3). Under these conditions, the electric and calcium waves could propagate  
356 through the vascular bundles (supported by GLR3.3;GLR3.6; Mousavi et al., 2013; Nguyen et  
357 al., 2018; Toyota et al., 2018; Shao et al., 2020), while the ROS wave could propagate through  
358 mesophyll cells (supported by RBOHD; Figures 1 and 3; Zandalinas et al., 2020b). This  
359 possibility suggests that the ROS wave can be spatially separated from the calcium and electric  
360 waves. Different stresses could therefore trigger different combinations of waves that could  
361 travel through different tissues and cell layers of the plant. During systemic responses to HL  
362 stress however the separation of systemic signals cannot occur (for reasons unknown at present),  
363 and the ROS wave must propagate together with the electric and calcium waves through the  
364 vascular bundles. Further studies are of course needed to address these intriguing possibilities.

365 Under all stresses studied here (HL, HS, wounding), RBOHD appeared to be required for  
366 systemic transcript accumulation (Figure 3), suggesting that even though GLRs were present and  
367 most likely functional in the *rbohD* mutant, they could not mediate their function to drive the  
368 expression of systemic transcript accumulation in the absence of the ROS wave. The ROS wave,  
369 even occurring at tissues other than the vascular bundles (*i.e.*, mesophyll cell layers; Figures 1  
370 and 2) could therefore be required to support other systemic signal propagation (such as electric  
371 and calcium waves) occurring at the vascular bundles during HS or wound responses. Although  
372 it is unknown at present how changes in ROS at the mesophyll cell layers impact electric and  
373 calcium signaling at the vascular bundles, one intriguing possibility is that different metabolites,  
374 ions, ROS, hormones, and/or pH changes, occurring at the mesophyll cell layers are  
375 diffused/transported to the vascular bundles, and these are needed to link the different waves  
376 (Fichman et al., 2020a; Fichman and Mittler, 2020). In this respect, it should be mentioned that  
377 changes in localized pH levels were recently linked to the triggering and propagation of electric  
378 and calcium waves in *Arabidopsis* (Shao et al., 2020). An alternative explanation could of course  
379 be that RBOHF at the vascular tissues replaces the function of RBOHD in linking the different  
380 waves during all stresses studied, and that the levels of RBOHF-produced ROS in the vascular  
381 bundles of *rbohD* plants complemented by *RBOHD* expressed under the control of a mesophyll-

382 specific promoter are too low to be detected by our assay. Further studies are of course needed to  
383 address these possibilities, as well as to resolve the different spatial and temporal relationships  
384 that could potentially exist between the different waves, signals and hormones involved in  
385 systemic signaling (*e.g.*, Kangasjärvi et al., 2009; Miller et al., 2009; Mittler et al., 2011;  
386 Dubiella et al., 2013; Gilroy et al., 2014; Evans et al., 2016; Gilroy et al., 2016; Choi et al., 2017;  
387 Fichman et al., 2020a; Fichman and Mittler, 2020).

388 Our findings that the ROS wave can propagate through multiple cell layers in response to  
389 different stresses could also partially explain how the integration of different systemic signals  
390 during a combination of HL and HS results in a stronger ROS wave signal (Zandalinas et al.,  
391 2020a). It is possible that during a combination of HL and HS applied to two different leaves of  
392 the same plant (HL&HS; Zandalinas et al., 2020a), the ROS wave initiated from the two  
393 different leaves propagates through all three cell types of the plant (mesophyll, xylem  
394 parenchyma and phloem, initiated by the local HS treatment, and xylem parenchyma and  
395 phloem, initiated by the local HL treatment). In contrast, during a combination of HL+HS  
396 applied to the same leaf, JA suppresses the ROS wave and the signal is lower (Zandalinas et al.,  
397 2020a). Our findings that restoring *RBOHD* expression in mesophyll cells did not result in a  
398 stronger systemic ROS signal during a HL&HS treatment (Figure 5), reveals that during HL&HS  
399 combination in *Arabidopsis* the ROS wave could indeed propagate through mesophyll cells  
400 (Figures 5 and 6 and Supplementary Figure S5). The ROS wave triggered by the HL treatment  
401 (propagating through xylem parenchyma and/or phloem cells) could therefore merge with the  
402 ROS wave triggered by the HS treatment (propagating through mesophyll and xylem  
403 parenchyma and/or phloem cells) to generate a stronger systemic ROS signal during HL&HS  
404 combination that is mediated through multiple cell layers (Figures 5 and 6 and Supplementary  
405 Figure S5). Because complementing *RBOHD* expression in the *rbohD* mutant using *RBOHD*  
406 expressed under the phloem-specific promoter was sufficient to restore the strong signal  
407 observed during HL&HS combination (Zandalinas et al., 2020a; Figure 5, Supplementary Figure  
408 S5), it is also possible that the stronger signal observed during HL&HS combination is simply  
409 the result of two different ROS wave signals merging together, regardless of the type of tissue  
410 supporting their transmission. Further studies are of course needed to dissect the mode of  
411 systemic signal integration through the different cell layers during stress combination.

412

## 413 MATERIALS AND METHODS

### 414 Plant material, growth conditions and stress treatments

415 *Arabidopsis thaliana* Col-0 (cv. Columbia-0), *rbohD* plants (Fichman et al., 2019) and two  
416 independent lines each of the different *rbohD* complemented plants (Zandalinas et al., 2020b)  
417 were grown in peat pellets (Jiffy-7, Jiffy, <http://www.jiffygroup.com/>) at 23°C under short day  
418 growth conditions (10-hour light/14-hour dark, 50  $\mu\text{mol m}^{-2} \text{s}^{-1}$ ). Wounding was achieved by  
419 puncturing a single leaf with 18 dressmaker pins (Singer, Murfreesboro, TN, USA) as described  
420 in (Fichman et al., 2019). Heat stress (HS) was induced by placing a heat block 2 cm underneath  
421 the treated leaf for 2 min, increasing the leaf temperature to 31-33°C (Zandalinas et al., 2020a).  
422 High light stress was applied by subjecting a single leaf to a light intensity of 1700  $\mu\text{mol m}^{-2} \text{s}^{-1}$   
423 for 2 min using a ColdVision fiber optic LED light source (Schott A20980, Southbridge, MA,  
424 USA) as described in (Devireddy et al., 2018; Zandalinas et al., 2019; Zandalinas et al., 2020a;  
425 Zandalinas et al., 2020b). The spectrum of this light stress treatment was shown in previous  
426 studies to contain all components required for triggering the systemic ROS signal through  
427 phytochrome B-mediated signaling (Devireddy et al., 2020b), as well as, when applied for more  
428 than 45 min, cause photosynthetic inhibition and light-induced cell death (Balfagón et al., 2019;  
429 Zandalinas et al., 2019; Zandalinas et al., 2020a; Zandalinas et al., 2020b). However, when  
430 applied for 2 min, this light stress treatment did not increase leaf temperature (Supplementary  
431 Table S1; Zandalinas et al., 2020a). Local and systemic leaf temperatures were measured under  
432 all conditions and treatment using an infrared camera (C2; FLIR Systems; Zandalinas et al.,  
433 2020a).

### 434 Measurements of ROS accumulation

435 To image whole-plant ROS levels, plants were fumigated with 50  $\mu\text{M}$   
436  $\text{H}_2\text{DCFDA}$  (excitation/emission 495 nm/517 nm; Millipore-Sigma, St. Louis, MO, USA) in  
437 50 mM phosphate buffer (pH 7.4) containing 0.01% Silwet L-77 (LEHLE seeds, Round Rock,  
438 TX, USA), using a portable mini nebulizer (Punasi Direct, Hong Kong, China) for 30 min as  
439 described previously (Fichman et al., 2019; Zandalinas et al., 2020a; Zandalinas et al., 2020b).  
440 Following  $\text{H}_2\text{DCFDA}$  application, local leaves were exposed to wounding, HL stress, HS, or HL

441 and HS applied to two different leaves located at opposite sides of the plant as described by  
442 Zandalinas et al., (2020a). Imaging of ROS accumulation in response to a local stress treatment  
443 was conducted with an IVIS Lumina S5 platform using Living Image 4.7.2 software  
444 (PerkinElmer) as described in (Fichman et al., 2019; Zandalinas et al., 2020a; Zandalinas et al.,  
445 2020b). All experiments were repeated at least three times each with 10 wild type, *rbohD* and the  
446 different complemented plants.

#### 447 **RT-qPCR analysis**

448 To analyze transcript expression by RT-qPCR, plants were subjected to a local treatment of  
449 wounding, 8-min HL or 8-min HS as described above. Local and systemic leaves were collected  
450 and immediately frozen in liquid nitrogen following the 8-min HL or HS treatments, or 30 min  
451 following wounding. Relative expression analysis by RT-qPCR was performed according to  
452 (Balfagón et al., 2019) by using the CFX Connect Real-Time PCR Detection System (Bio-Rad)  
453 and gene-specific primers (Supplementary Table S2; Primer efficiency range of 0.99-1.04). All  
454 experiments were repeated at least three times each with at least 5 wild type, *rbohD* and the  
455 different *rbohD* complemented plants.

#### 456 **Heat stress acclimation assay**

457 For heat stress acclimation, a single leaf was pre-treated for 15 min at 31-33°C by placing a heat  
458 block 2 cm underneath the treated leaf (Zandalinas et al., 2020a). Plants were then incubated for  
459 45 minutes under controlled conditions. Following the recovery period, a systemic leaf of pre-  
460 treated and untreated plants was dipped in a 42°C (or 23°C as control) water bath for 60 min and  
461 allowed to recover under controlled growth conditions. Systemic leaves were sampled 6 days  
462 after the water bath heat stress treatment for chlorophyll measurements, as previously described  
463 (Zandalinas et al., 2020a; Zandalinas et al., 2020c). For HL-induced SAA, a single leaf was pre-  
464 treated for 15 min with a light intensity of 1700  $\mu\text{mol m}^{-2} \text{sec}^{-1}$  using a ColdVision fiber optic  
465 LED light source (Schott A20980, Southbridge, MA, USA). Plants were then incubated for 45  
466 minutes under controlled conditions. Following the recovery period, a systemic leaf was exposed  
467 to a light intensity of 1700  $\mu\text{mol m}^{-2} \text{sec}^{-1}$  for 45 minutes. Control systemic leaves (untreated)  
468 and systemic leaves of plants that were pretreated with HL stress, as described above (SAA),  
469 were then analyzed for electrolyte leakage as previously described (Zandalinas et al., 2019;



470 Zandalinas et al., 2020a; Zandalinas et al., 2020b). Acclimation assays were repeated at least 3  
471 times with 10 plants per repeat.

## 472 **GFP imaging**

473 Localization of RBOHD-GFP in leaves of mature (4-5-week-old) *rbohD* plants complemented  
474 with the RBOHD-GFP protein driven by its native or CAB3 promoter was performed using a  
475 TCS SP8 (Leica) multiphoton confocal microscope (Buffalo Grove, IL, USA) as described in  
476 Zandalinas et al., 2020b.

## 477 **Statistical analysis**

478 Results are presented as the mean  $\pm$  SD. Statistical analyses were performed by a two-tailed  
479 Student's t-test (asterisks denote statistical significance at  $p < 0.05$  with respect to controls).

480

## 481 **SUPPLEMENTAL DATA**

482 **Supplementary Table S1.** FLIR camera measurements showing the surface temperature of  
483 treated (local) and systemic leaves for each stress treatment (C2, FLIR systems AB).  
484 *Abbreviations used:* CT, control; HL, high light; HS, heat stress.

485 **Supplementary Table S2.** Transcript-specific primers used for relative expression analysis by  
486 RT-qPCR.

487 **Supplementary Figure S1.** Representative confocal images of RBOHD-GFP fusion protein  
488 expression in mature leaves of transgenic *rbohD* mutants. The RBOHD-GFP protein was  
489 expressed in the *rbohD* mutant background under the control of the native RbohD or the CAB3  
490 promoters. *Abbreviations used:* RBOHD, respiratory burst oxidase homolog D; CAB,  
491 chlorophyll A/B binding protein (Scale bar = 20  $\mu$ m).

492 **Supplementary Figure S2.** Linear regression analysis conducted using scatter plots of  
493 continuous ROS measurements in local and systemic leaves of wild type, *rbohD* and the  
494 different complemented lines over the entire course of each experiment (0 to 30 min). Best-fit  
495 regression lines (in black) and the slope of each signal progression are shown. *Abbreviations*  
496 *used:* HL, high light; HS, heat stress; RBOHD, respiratory burst oxidase homolog D; CER,

497 eceriferum; CAB, chlorophyll A/B binding protein; SCR, scarecrow; XCP, xylem cysteine  
498 peptidase; Sultr, sulfate transporter; TRE, total radiant efficiency; W, wounding.

499 **Supplementary Figure S3.** Complementation of light (HL) stress-induced local and systemic  
500 ROS signaling in the *rbohD* mutant with *RBOHD* driven by different tissue-specific promoters.  
501 Representative time-lapse images of whole-plant ROS levels in wild type, *rbohD* and the  
502 different *rbohD* complemented *Arabidopsis thaliana* plants subjected to a local HL-stress  
503 treatment (red circles), are shown on left; representative line graphs showing continuous  
504 measurements of ROS levels in local and systemic leaves of wild type, *rbohD* and two  
505 independent homozygous complemented lines (#1 and #2), over the entire course of the  
506 experiment (0 to 30 min) are shown in the middle (ROIs for some of them are indicated with  
507 blue boxes); and statistical analysis of ROS levels in local and systemic leaves at 0 and 30 min is  
508 shown on right (Student t-test, SD, N=10, \*p < 0.05). All experiments were repeated at least 3  
509 times with similar results. Scale bar indicates 1 cm. *Abbreviations used:* HL, high light; RBOHD,  
510 respiratory burst oxidase homolog D; CER, eceriferum; CAB, chlorophyll A/B binding protein;  
511 SCR, scarecrow; XCP, xylem cysteine peptidase; ROI, region of interest; Sultr, sulfate  
512 transporter; TRE, total radiant efficiency. The experiments shown were conducted in parallel to  
513 the experiments shown in Figures 1 and 2 and are a repeat of the study reported previously  
514 (Zandalinas et al., 2020b), with similar results.

515 **Supplementary Figure S4.** Complementation of light stress (HL)-induced SAA in the *rbohD*  
516 mutant with *RBOHD* driven by different tissue-specific promoters. Light stress-induced systemic  
517 leaf cell injury (measured as electrolyte leakage) of wild type, *rbohD* and the different *rbohD*-  
518 complemented *Arabidopsis thaliana* plants is shown. Systemic leaves were either untreated and  
519 unstressed (Control) or subjected to a systemic light stress following a local pretreatment of a  
520 local leaf with light stress (SAA). Ten different plants each from two independent complemented  
521 lines for each construct were subjected to light stress and cell injury was determined by  
522 measuring electrolyte leakage from systemic leaves. Student t-test, SD, N=10, \*p < 0.05.  
523 *Abbreviations used:* HL, high light; RBOHD, respiratory burst oxidase homolog D; CER,  
524 eceriferum; CAB, chlorophyll A/B binding protein; SCR, scarecrow; XCP, xylem cysteine  
525 peptidase; Sultr, sulfate transporter; EL, electrolyte leakage; SAA, systemic acquired  
526 acclimation. The experiments shown were conducted in parallel to the experiments shown in

527 Figure 4 and are a repeat of the study reported previously (Zandalinas et al., 2020b), with similar  
528 results.

529 **Supplementary Figure S5.** Complementation of light (HL)- and heat (HS)-induced local and  
530 systemic ROS signaling in the *rbohD* mutant with *RBOHD* driven by the phloem- or mesophyll  
531 tissue-specific promoters, during stress combination. Representative images of whole-plant ROS  
532 levels in *rbohD* and *rbohD*-complemented *Arabidopsis thaliana* plants 20 min following a local  
533 light (HL)- or heat (HS)- treatments, or a combination of light- and heat-stress treatments applied  
534 to two leaves of the same plant (HL&HS; Zandalinas et al., 2020a; red circles) are shown. All  
535 experiments were repeated at least 3 times with similar results. Scale bar indicates 1 cm.  
536 *Abbreviations used:* HL, high light; HS, heat stress; RBOHD, respiratory burst oxidase homolog  
537 D; CAB, chlorophyll A/B binding protein; Sultr, sulfate transporter.

538 **FIGURE LEGENDS**

539 **Figure 1.** Complementation of wound-induced local and systemic ROS signaling in the *rbohD*  
540 mutant with *RBOHD* driven by different tissue-specific promoters. Representative time-lapse  
541 images of whole-plant ROS levels in wild type, *rbohD* and the different *rbohD*-complemented  
542 *Arabidopsis thaliana* plants subjected to a local wound treatment (red circles), are shown on left;  
543 representative line graphs showing continuous measurements of ROS levels in local and  
544 systemic leaves of wild type, *rbohD*, and two independent homozygous complemented lines (#1  
545 and #2), over the entire course of the experiment (0 to 30 min) are shown in the middle (ROIs for  
546 some of them are indicated with blue boxes); and statistical analysis of ROS levels in local and  
547 systemic leaves at 0 and 30 min is shown on right (Student t-test, SD, N=10, \*p < 0.05). All  
548 experiments were repeated at least 3 times with similar results. Scale bar indicates 1 cm.  
549 *Abbreviations used:* RBOHD, respiratory burst oxidase homolog D; CER, eceriferum; CAB,  
550 chlorophyll A/B binding protein; SCR, scarecrow; XCP, xylem cysteine peptidase; ROI, region  
551 of interest; Sultr, sulfate transporter; TRE, total radiant efficiency.

552 **Figure 2.** Complementation of heat stress-induced local and systemic ROS signaling in the  
553 *rbohD* mutant with *RBOHD* driven by different tissue-specific promoters. Representative time-  
554 lapse images of whole-plant ROS levels in wild type, *rbohD* and the different *rbohD*-  
555 complemented *Arabidopsis thaliana* plants subjected to a local heat stress treatment (red circles),  
556 are shown on left; representative line graphs showing continuous measurements of ROS levels in  
557 local and systemic leaves of wild type, *rbohD* and two independent homozygous complemented  
558 lines (#1 and #2), over the entire course of the experiment (0 to 30 min) are shown in the middle  
559 (ROIs for some of them are indicated with blue boxes); and statistical analysis of ROS levels in  
560 local and systemic leaves at 0 and 30 min is shown on right (Student t-test, SD, N=10, \*p <  
561 0.05). All experiments were repeated at least 3 times with similar results. Scale bar indicates 1  
562 cm. *Abbreviations used:* RBOHD, respiratory burst oxidase homolog D; CER, eceriferum; CAB,  
563 chlorophyll A/B binding protein; SCR, scarecrow; XCP, xylem cysteine peptidase; ROI, region  
564 of interest; Sultr, sulfate transporter; TRE, total radiant efficiency.

565 **Figure 3.** Local- and systemic stress-induced transcript expression in wild type, *rbohD*, and the  
566 *rbohD* mutant complemented with *RBOHD* driven by different tissue-specific promoters. (A)  
567 Local and systemic steady-state levels of *JAZ5* (AT1G17380) and *JAZ7* (AT2G34600)

568 transcripts in wild type, *rbohD*, and the different *rbohD*-complemented *Arabidopsis thaliana*  
569 plants subjected to a local wound treatment. (B) Local and systemic steady-state levels of *Rap2.4*  
570 (AT1G78080) and *ERF2* (AT5G47220) transcripts in wild type, *rbohD*, and the different *rbohD*-  
571 complemented *Arabidopsis* plants subjected to a local heat-stress treatment. (C) Local and  
572 systemic steady-state levels of *MYB30* (AT3G28910) and *ZHD5* (AT1G75240) transcripts in  
573 wild type, *rbohD*, and the different *rbohD*-complemented *Arabidopsis* plants subjected to a local  
574 high light-stress treatment. Student t-test, SD, N=3, \*p < 0.05. *Abbreviations used*: RBOHD,  
575 respiratory burst oxidase homolog D; CER, eceriferum; CAB, chlorophyll A/B binding protein;  
576 SCR, scarecrow; XCP, xylem cysteine peptidase; Sultr, sulfate transporter; JAZ, jasmonate-zim-  
577 domain protein; ERF, ethylene response factor; ZHD, zinc-finger homeodomain.

578 **Figure 4.** Complementation of heat stress-induced SAA in the *rbohD* mutant with *RBOHD*  
579 driven by different tissue-specific promoters. Heat stress-induced changes in systemic leaf  
580 chlorophyll content of wild type, *rbohD* and the different *rbohD*-complemented *Arabidopsis*  
581 *thaliana* plants are shown. Systemic leaves were obtained from plants that were either untreated  
582 and unstressed (Control), untreated at their local leaves and subjected to a systemic heat-stress  
583 treatment (No pretreatment), or subjected to a local heat stress pre-treatment before being  
584 subjected to a systemic heat stress treatment (Pretreatment). SAA is evident when the systemic  
585 leaf of a pre-treated plant does not show a loss of chlorophyll content following a systemic heat  
586 stress treatment. Ten different plants each from two independent complemented lines for each  
587 construct were subjected to the SAA heat stress assay and chlorophyll content was measured in  
588 systemic leaves. Student t-test, SD, N=10, \*p < 0.05. *Abbreviations used*: RBOHD, respiratory  
589 burst oxidase homolog D; CER, eceriferum; CAB, chlorophyll A/B binding protein; SCR,  
590 scarecrow; XCP, xylem cysteine peptidase; Sultr, sulfate transporter.

591 **Figure 5.** Complementation of light- and heat-induced local and systemic ROS signaling in the  
592 *rbohD* mutant with *RBOHD* driven by the phloem- or mesophyll tissue-specific promoters,  
593 during stress combination. Representative time-lapse images of whole-plant ROS levels of wild  
594 type, *rbohD* and *rbohD*-complemented *Arabidopsis thaliana* plants subjected to a local light  
595 (HL) and heat stress (HS) treatments, simultaneously applied to two leaves of the same plant (red  
596 circles; HL&HS; Zandalinas et al., 2020a) are shown on left (ROIs for some of them are  
597 indicated with blue boxes), and statistical analysis of ROS levels in systemic leaves of treated

598 plants at 0, 10, 15 and 20 min is shown on right (Student t-test, SD, N=10, \*p < 0.05). All  
599 experiments were repeated at least 3 times with similar results. Scale bar indicates 1 cm.  
600 *Abbreviations used:* HL, high light; HS, heat stress; RBOHD, respiratory burst oxidase homolog  
601 D; CAB, chlorophyll A/B binding protein; Sultr, sulfate transporter.

602 **Figure 6.** A model showing that when light stress is applied to a local leaf, the ROS wave is  
603 mediated through vascular bundles. In contrast, when heat stress or wounding are applied to a  
604 local leaf, both vascular and mesophyll cells can mediate the ROS wave. *Abbreviations used:*  
605 HL, high light; HS, heat stress; W, wounding; ROS, reactive oxygen species.

606

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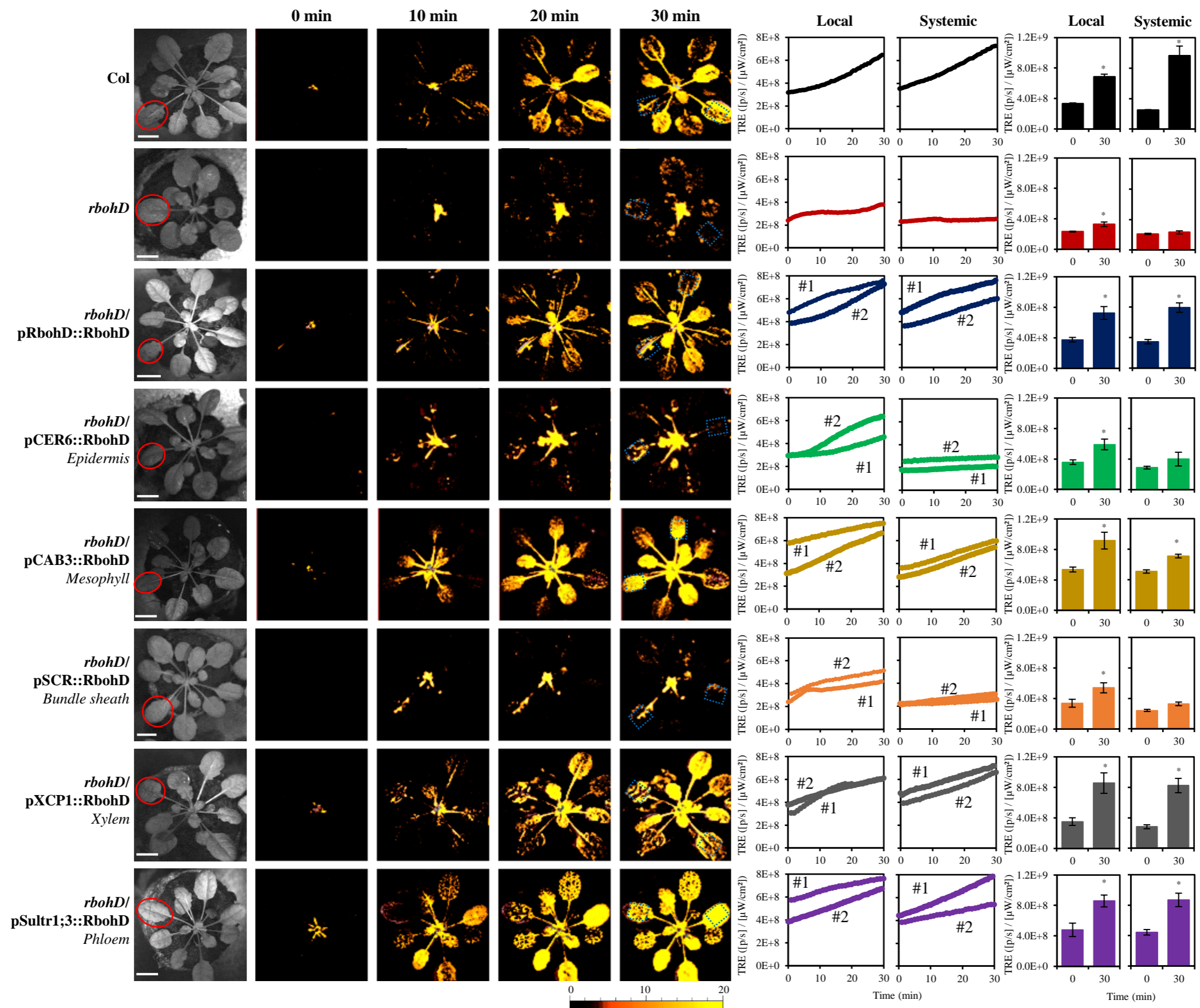
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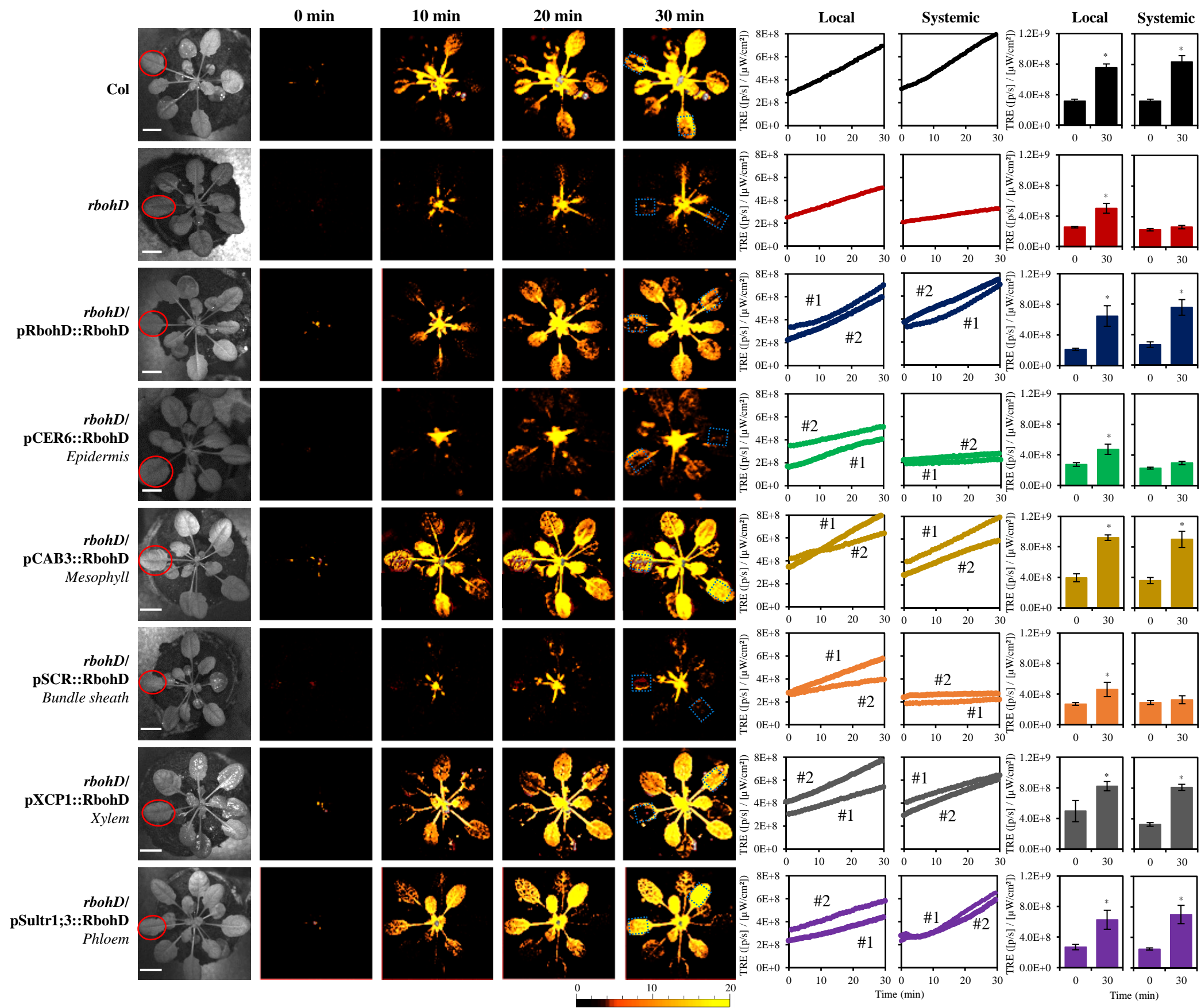
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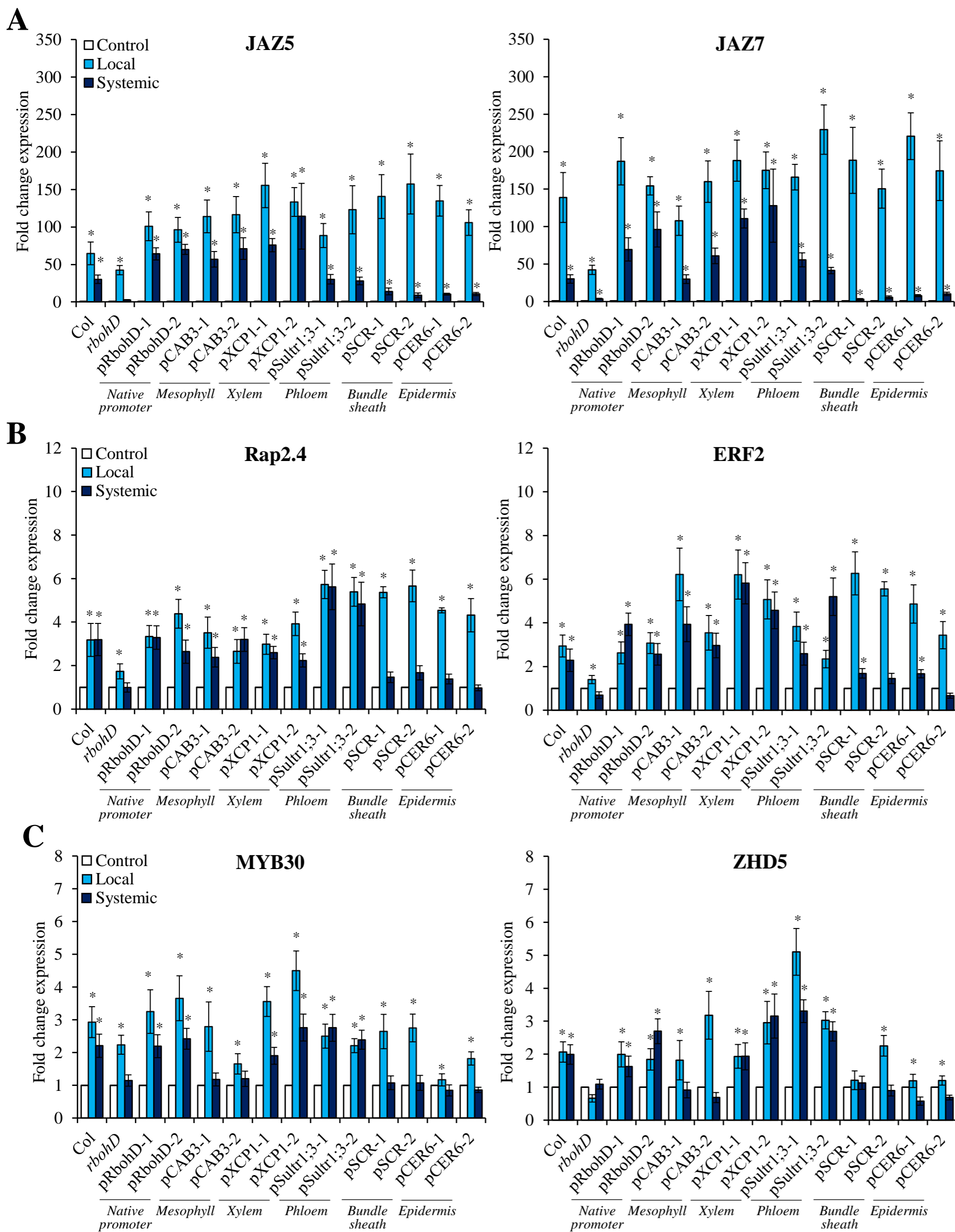
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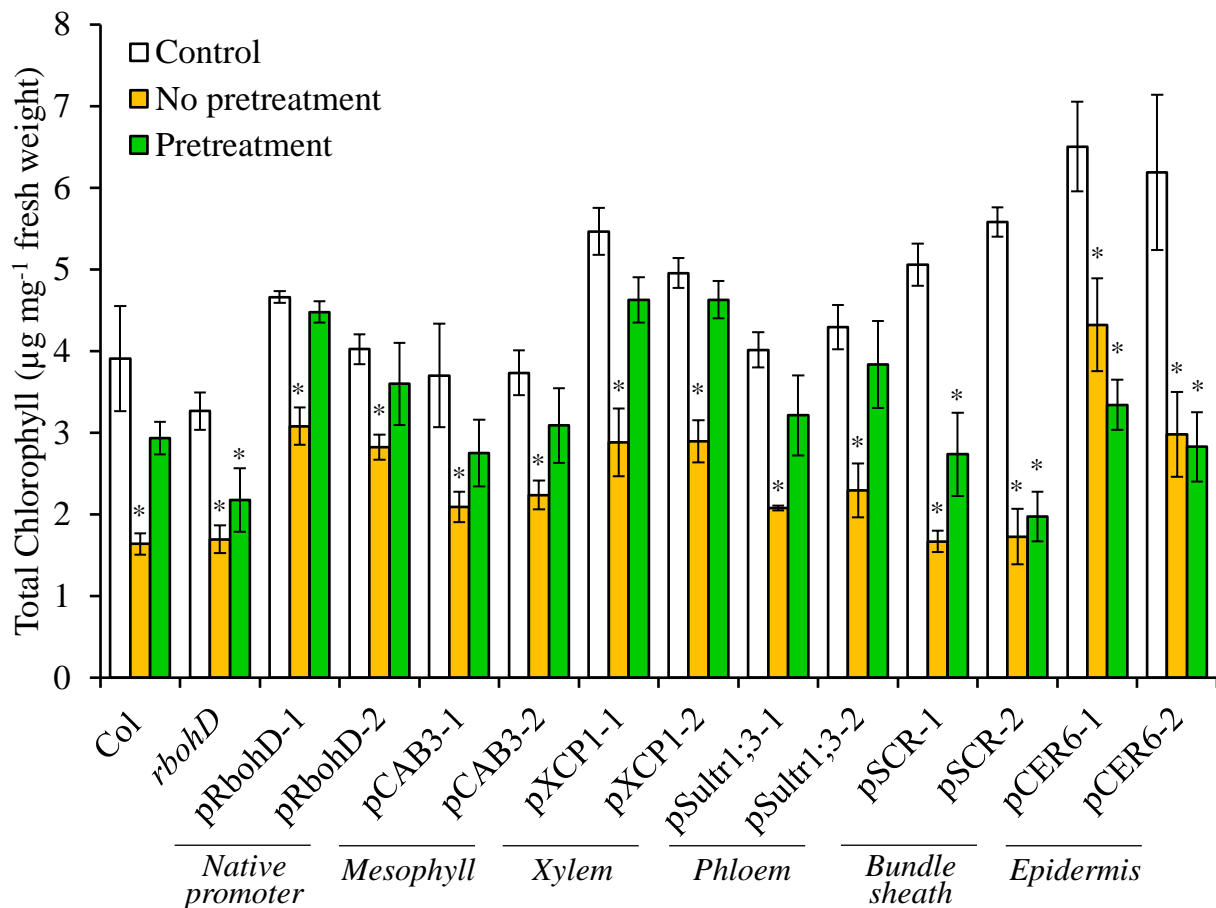
**Figure 1.** Complementation of wound-induced local and systemic ROS signaling in the *rbohD* mutant with *RBOHD* driven by different tissue-specific promoters. Representative time-lapse images of whole-plant ROS levels in wild type, *rbohD* and the different *rbohD*-complemented *Arabidopsis thaliana* plants subjected to a local wound treatment (red circles), are shown on left; representative line graphs showing continuous measurements of ROS levels in local and systemic leaves of wild type, *rbohD*, and two independent homozygous complemented lines (#1 and #2), over the entire course of the experiment (0 to 30 min) are shown in the middle (ROIs for some of them are indicated with blue boxes); and statistical analysis of ROS levels in local and systemic leaves at 0 and 30 min is shown on right (Student t-test, SD, N=10, \**p* < 0.05). All experiments were repeated at least 3 times with similar results. Scale bar indicates 1 cm. *Abbreviations used:* RBOHD, respiratory burst oxidase homolog D; CER, eceriferum; CAB, chlorophyll A/B binding protein; SCR, scarecrow; XCP, xylem cysteine peptidase; ROI, region of interest; Sultr, sulfate transporter; TRE, total radiant efficiency.



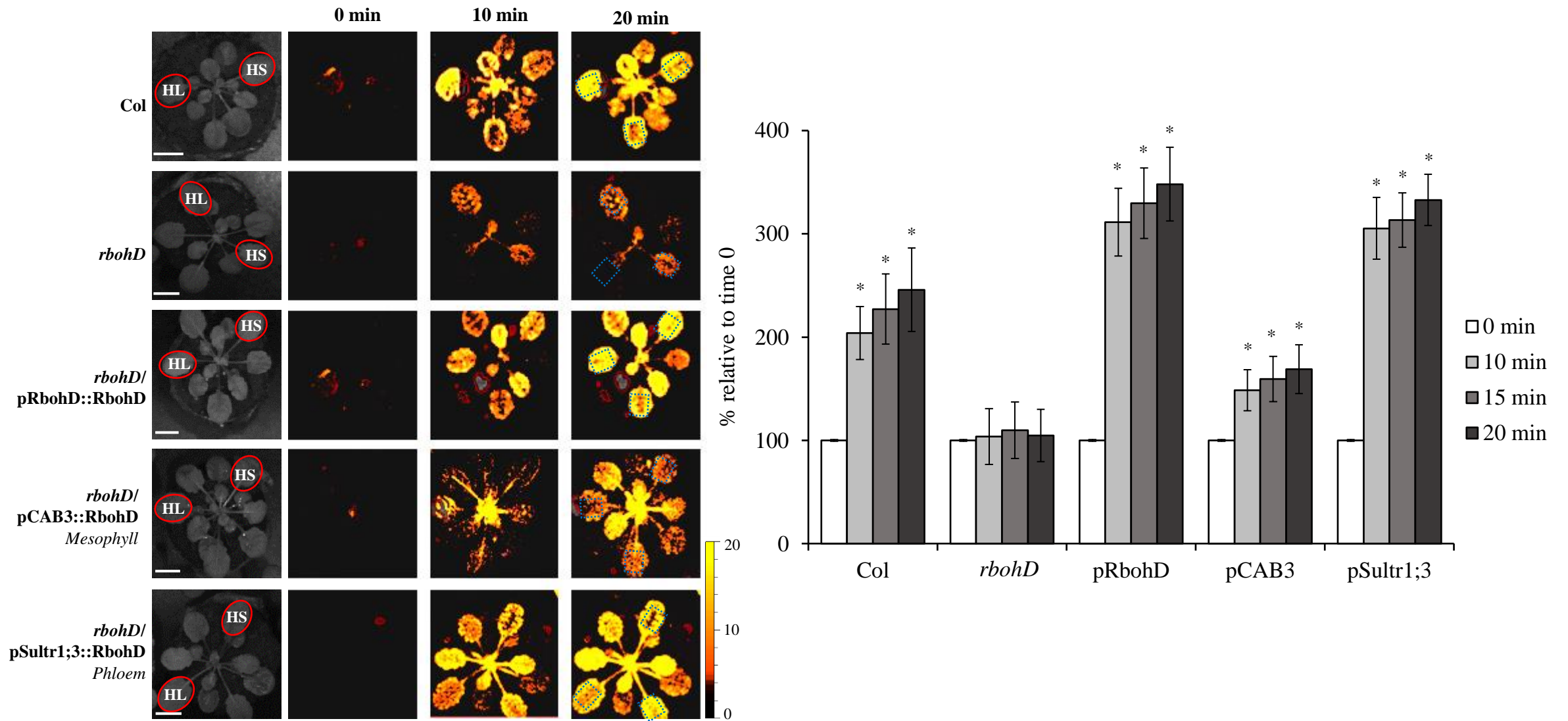
**Figure 2.** Complementation of heat stress-induced local and systemic ROS signaling in the *rbohD* mutant with *RBOHD* driven by different tissue-specific promoters. Representative time-lapse images of whole-plant ROS levels in wild type, *rbohD* and the different *rbohD*-complemented *Arabidopsis thaliana* plants subjected to a local heat stress treatment (red circles), are shown on left; representative line graphs showing continuous measurements of ROS levels in local and systemic leaves of wild type, *rbohD* and two independent homozygous complemented lines (#1 and #2), over the entire course of the experiment (0 to 30 min) are shown in the middle (ROIs for some of them are indicated with blue boxes); and statistical analysis of ROS levels in local and systemic leaves at 0 and 30 min is shown on right (Student t-test, SD, N=10, \*p < 0.05). All experiments were repeated at least 3 times with similar results. Scale bar indicates 1 cm. *Abbreviations used:* RBOHD, respiratory burst oxidase homolog D; CER, eceriferum; CAB, chlorophyll A/B binding protein; SCR, scarecrow; XCP, xylem cysteine peptidase; ROI, region of interest; Sultr, sulfate transporter; TRE, total radiant efficiency.



**Figure 3.** Local and systemic stress-induced transcript expression in wild type, *rbohD*, and the *rbohD* mutant complemented with *RBOHD* driven by different tissue-specific promoters. (A) Local and systemic steady-state levels of *JAZ5* (AT1G17380) and *JAZ7* (AT2G34600) transcripts in wild type, *rbohD*, and the different *rbohD*-complemented *Arabidopsis thaliana* plants subjected to a local wound treatment. (B) Local and systemic steady-state levels of *Rap2.4* (AT1G78080) and *ERF2* (AT5G47220) transcripts in wild type, *rbohD*, and the different *rbohD*-complemented *Arabidopsis* plants subjected to a local heat-stress treatment. (C) Local and systemic steady-state levels of *MYB30* (AT3G28910) and *ZHD5* (AT1G75240) transcripts in wild type, *rbohD*, and the different *rbohD*-complemented *Arabidopsis* plants subjected to a local high light-stress treatment. Student t-test, SD, N=3, \* $p < 0.05$ . Abbreviations used: RBOHD, respiratory burst oxidase homolog D; CER, eceriferum; CAB, chlorophyll A/B binding protein; SCR, scarecrow; XCP, xylem cysteine peptidase; Sultr, sulfate transporter; JAZ, jasmonate-zim-domain protein; ERF, ethylene response factor; ZHD, zinc-finger homeodomain.

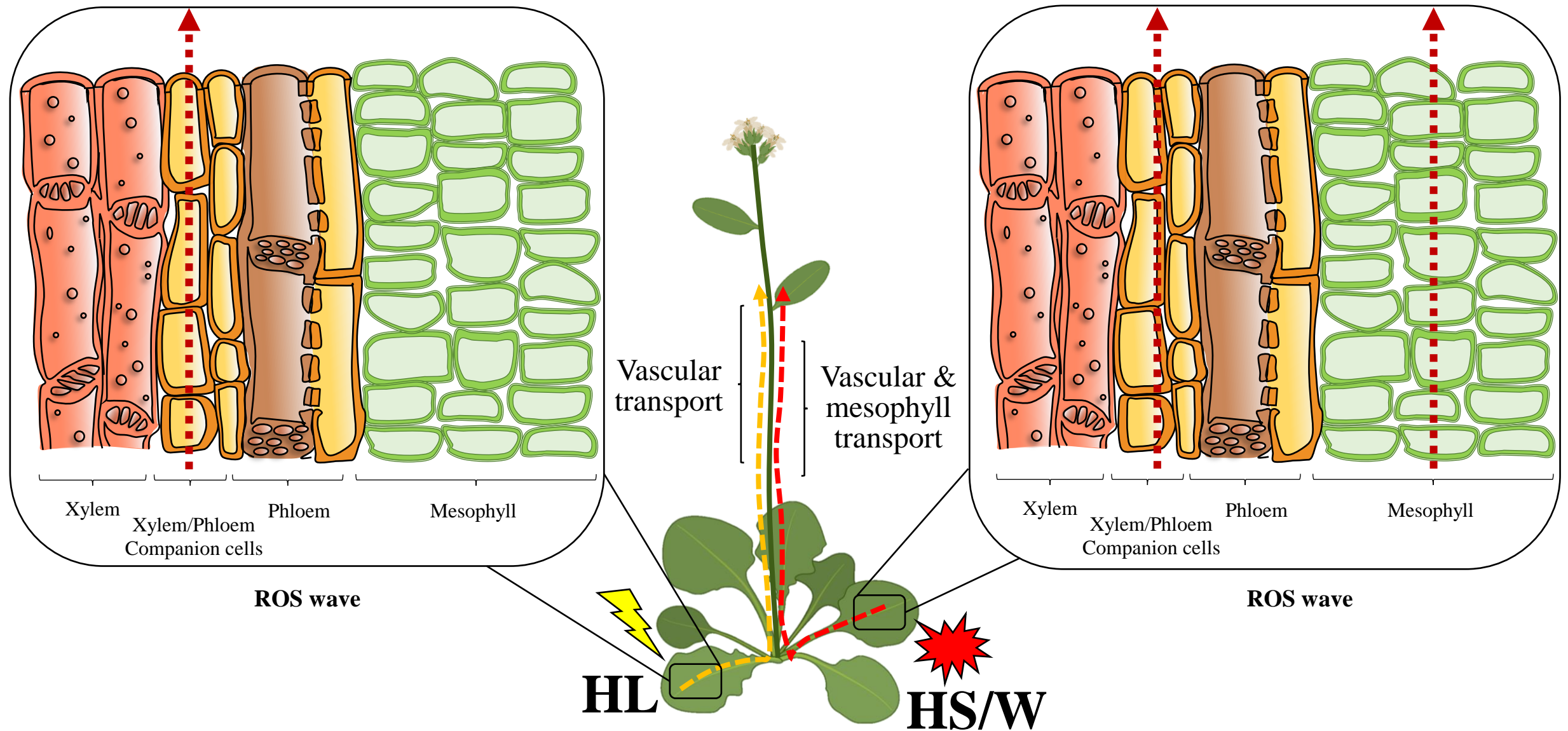


**Figure 4.** Complementation of heat stress-induced SAA in the *rbohD* mutant with *RBOHD* driven by different tissue-specific promoters. Heat stress-induced changes in systemic leaf chlorophyll content of wild type, *rbohD* and the different *rbohD*-complemented *Arabidopsis thaliana* plants are shown. Systemic leaves were obtained from plants that were either untreated and unstressed (Control), untreated at their local leaves and subjected to a systemic heat-stress treatment (No pretreatment), or subjected to a local heat stress pre-treatment before being subjected to a systemic heat stress treatment (Pretreatment). SAA is evident when the systemic leaf of a pre-treated plant does not show a loss of chlorophyll content following a systemic heat stress treatment. Ten different plants each from two independent complemented lines for each construct were subjected to the SAA heat stress assay and chlorophyll content was measured in systemic leaves. Student t-test, SD, N=10, \* $p < 0.05$ . *Abbreviations used:* RBOHD, respiratory burst oxidase homolog D; CER, eceriferum; CAB, chlorophyll A/B binding protein; SCR, scarecrow; XCP, xylem cysteine peptidase; Sultr, sulfate transporter.



**Figure 5.** Complementation of light- and heat-induced local and systemic ROS signaling in the *rbohD* mutant with *RBOHD* driven by the phloem- or mesophyll tissue-specific promoters, during stress combination. Representative time-lapse images of whole-plant ROS levels of wild type, *rbohD* and *rbohD*-complemented *Arabidopsis thaliana* plants subjected to a local light (HL) and heat stress (HS) treatments, simultaneously applied to two leaves of the same plant (red circles; HL&HS; Zandalinas et al., 2020a) are shown on left (ROIs for some of them are indicated with blue boxes), and statistical analysis of ROS levels in systemic leaves of treated plants at 0, 10, 15 and 20 min is shown on right (Student t-test, SD, N=10, \* $p < 0.05$ ). All experiments were repeated at least 3 times with similar results. Scale bar indicates 1 cm. *Abbreviations used:* HL, high light; HS, heat stress; RBOHD, respiratory burst oxidase homolog D; CAB, chlorophyll A/B binding protein; Sultr, sulfate transporter.





**Figure 6.** A model showing that when light stress is applied to a local leaf, the ROS wave is mediated through vascular bundles. In contrast, when heat stress or wounding are applied to a local leaf, both vascular and mesophyll cells can mediate the ROS wave. *Abbreviations used:* HL, high light; HS, heat stress; W, wounding; ROS, reactive oxygen species.

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