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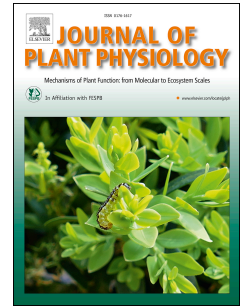


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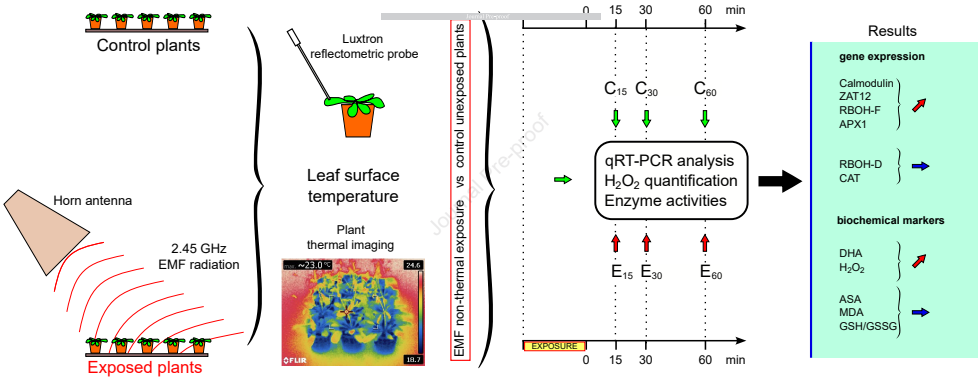
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2 *thaliana* metabolism.

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24 **Abstract**

25

26

27 Numerous studies report different types of responses following exposure of plants to high  
28 frequency electromagnetic fields (HF-EMF). While this phenomenon is related to tissue heating  
29 in animals, the situation is much less straightforward in plants where metabolic changes seem  
30 to occur without tissue temperature increase. We have set up an exposure system allowing  
31 reliable measurements of tissue heating (using a reflectometric probe and thermal imaging)  
32 after a long exposure (30 min) to an electromagnetic field of 2.45 GHz transmitted through a  
33 horn antenna (about  $100 \text{ V.m}^{-1}$  at the plant level). We did not observe any heating of the  
34 tissues, but we detected rapid increases (60 min) in the accumulation of transcripts of stress-  
35 related genes (*TCH1* and *ZAT12* transcription factor) or involved in ROS metabolism (*RBOHF*  
36 and *APX1*). At the same time, the amounts of hydrogen peroxide and dehydroascorbic acid  
37 increased while glutathione (reduced and oxidized forms), ascorbic acid, and lipid peroxidation  
38 remained stable. Therefore, our results unambiguously show that molecular and biochemical  
39 responses occur rapidly (within 60min) in plants after exposure to an electromagnetic field, in  
40 absence of tissue heating.

41

42 **Keywords** : electromagnetic exposure, non-thermal, stress-marker genes, hydrogen  
43 peroxide

44

45 **Abbreviations:**

46 HF-EMF: High Frequency-Electromagnetic Field

47 SAR: Specific Absorption Rate

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## 51 **Introduction**

52 In nature, plants are increasingly subjected to various kind of high frequency electromagnetic  
53 fields (HF-EMF) due to the densification of base station networks that comply with the exposure  
54 limits defined by the standards (ICNIRP, 2020; Israel et al., 2013). The question of whether  
55 high-frequency electromagnetic fields are likely to induce changes in plant metabolism and/or  
56 development remains however largely open, although many studies tend to show that they do  
57 (Kaur et al., 2021; Vian et al., 2016). Biological effects of electromagnetic field are generally  
58 interpreted in term of specific absorption rate (SAR) that relies on tissue heating after exposure  
59 to HF-EMF. No clear biological changes have been proven to occur in animal or human cells  
60 after EMF exposure in the absence of thermal effect (Habauzit et al., 2014; Roux et al., 2010).  
61 However, some studies clearly suggested that it could be different in plants in which various  
62 kind of biological responses have been detected after exposure to low amplitude, non-thermal  
63 EMF. Indeed, Halgamuge et al. (2015) have shown that soybean seedlings exposed to various  
64 levels of 900 MHz HF-EMF (SAR of  $4.8 \cdot 10^{-7}$  to  $20 \cdot 10^{-3} \text{ Wkg}^{-1}$ ) for durations ranging from 2h to  
65 5 days induced changes (mainly growth inhibition) of the plant root and hypocotyl. Similarly,  
66 4h exposure to HF-EMF (1800 MHz, SAR of  $1.69 \cdot 10^{-1} \text{ Wkg}^{-1}$ ) affected root development,  
67 coleoptile length and total chlorophyll content in maize (Kumar et al., 2016). Short exposure of  
68 rose buds to low amplitude HF-EMF (900 MHz, SAR of  $7.2 \cdot 10^{-4} \text{ W kg}^{-1}$ ) in a mode stirred  
69 reverberation chamber (MSRC) did not change the length of the axis produced after bud  
70 outgrowth, but the post-formed axillary buds produced 45% shorter axis, suggesting that  
71 growth changes may occur in a delayed manner (Grémiaux et al., 2016). Recently, Upadhyaya  
72 et al. (2022) noted after 12-120h exposure to HF-EMF (1800 MHz , SAR of  $3.16 \cdot 10^{-2} \text{ W kg}^{-1}$ )  
73 a decrease of seed germination, a reduced growth of plantlets, a reduction of leaf size and an  
74 increase in  $\text{H}_2\text{O}_2$  production. Environmental sensing and signaling in plants are largely  
75 dependent on reactive oxygen species (ROS) as signaling molecules, particularly  $\text{H}_2\text{O}_2$  which  
76 is the most stable and transportable ROS, involved in subcellular signaling as well as in cell-  
77 to-cell signaling (Mittler et al., 2022). Singh et al. (2012) observed reduced growth of mung  
78 bean seedlings hypocotyl and roots after 2h low amplitude HF-EMF exposure, along with a

79 4.5-fold elevation of H<sub>2</sub>O<sub>2</sub> content and to concomitant increase of the activities of several  
80 antioxidant enzymes: superoxide dismutase (4-fold), catalase (3.3-fold), ascorbate peroxidase  
81 (2.5-fold), guaiacol peroxidase (2-folds) and glutathione reductase (5-fold). Similar finding was  
82 reported by Chandel et al. (2017) in *Allium cepa* roots after 4h exposure to low level HF-EMF  
83 (2100 MHz, SAR of 2.82 10<sup>-1</sup> W kg<sup>-1</sup>): H<sub>2</sub>O<sub>2</sub> content was increased by 5-fold, while superoxide  
84 dismutase and catalase activities were up-regulated 2.4-fold. Gene expression was also  
85 affected after exposing plants to low level (non-thermal) HF-EMF: tomato plants exposed to  
86 10 min at 900 MHz (5 Vm<sup>-1</sup>) displayed rapid and transient changes in the expression of several  
87 stress-related genes, namely calmodulin, calcium-dependent protein kinase, *bZIP*  
88 transcription factor, proteinase inhibitor (*PIN2*) and Chloroplast mRNA Binding Protein (*CMBP*,  
89 Roux et al., 2006; Roux et al., 2008; Vian et al., 2006). These results were partially and  
90 independently replicated (Rammal et al., 2014), although these authors used a much less  
91 sophisticated exposure system and a different exposure condition (10 days, 1250 MHz, 6 Vm<sup>-1</sup>  
92 ). These changes occurred in a systemic way, since an exposure of a single leaf to HF-EMF  
93 triggered changes in gene expression not only locally (*i.e.* at the site where the exposure to  
94 HF-EMF was performed), but also at distance in the terminal leaf that was shielded from HF-  
95 EMF (Beaubois et al., 2007). More recently, Kundu et al. (2021a) showed that exposure of rice  
96 plants over a long period of time (6h per day for 32 days, 2.75 mW m<sup>-2</sup>) resulted in a significant  
97 increase in calmodulin, calcium dependent protein kinase and phytochrome C genes  
98 expression. Furthermore, the same research group (Kundu et al., 2021b) demonstrated in  
99 40 days-old 'Swarnaprabha' rice plants that a single HF-EMF irradiation (2 h 30 min,  
100 1837.50 MHz, 2.75 mWm<sup>-2</sup>) increased the expression level of some stress and light-signaling  
101 related genes: calmodulin (2.5-fold), *bZIP* transcription factor (2.27-fold) and phytochrome B  
102 and C (3.98- and 5.87-fold, respectively). A mechanism that integrates the rapid responses of  
103 plants to exposure to very high frequency electromagnetic waves has recently been proposed  
104 (Kaur et al., 2021).

105 In the present work, we used *Arabidopsis thaliana* as a plant model to assess whether a short  
106 (30 min), non-thermal exposure to HF-EMF could trigger rapid changes in plant metabolism.  
107 We used a horn antenna that delivers an electromagnetic signal with a fixed angle of incidence  
108 and polarization to get similar exposure configuration to those used in the vast majority of the  
109 works, in preference to the mode-stirring reverberation chamber that we used in our previous  
110 work (Vian et al., 2006; Roux et al., 2006), which allows to create a homogeneous and isotropic  
111 electromagnetic environment, but which remains an equipment that is seldomly used for  
112 bioelectromagnetic studies. We paid particular attention to ensure that the exposure (30 min,  
113 2450 MHz) did not induce thermal effects in the plants and investigate the possible resulting  
114 effects of this non-thermal exposure in terms of rapid (within one hour) changes in gene  
115 expression and biochemical pathways. Using these carefully controlled experimental  
116 conditions, we provide evidence that exposure to non-thermal HF-EMF was perceived by  
117 plants and induced metabolic changes in the rosette of *Arabidopsis thaliana*.

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133 **Materials and Methods**134 *Plant material and culture*

135 *Arabidopsis* (*Arabidopsis thaliana* ecotype Col-0) were grown in controlled environmental  
136 conditions in culture chambers (Binder KBW720) under 150-160  $\mu\text{mol.m}^{-2}.\text{s}^{-1}$  (provided by  
137 OSRAM Fluora L18W/77 fluorescent tubes),  $21\pm 1^\circ\text{C}$  at a 16h light period in an adapted  
138 substrate for young plants in module trays (TraySubstrat U44-551, Klasmann-Deilmann,  
139 Bourgoin Jallieu, France). The plants were watered by sub-irrigation with a nutritive solution  
140 and grown for 4 weeks to obtain rosettes with a diameter of 4-5 cm.

141 *Plant exposure to electromagnetic field and material sampling*

142 The experiments were set-up in a shielded environment (metal-walled room) to avoid possible  
143 interaction with the external (environmental) electromagnetic background. A signal synthesizer  
144 (SMB100A, Rohde & Schwarz, Munich, Germany) and modulator (AFG3021B, Tektronix,  
145 Beaverton, USA) produced a signal, amplified to its nominal amplitude by a high frequency  
146 amplifier (ASO104-30/17, Milmega, Ryde, United Kingdom) and frequency modulated by a  
147 100 kHz square signal. Thus, the amplitude of the signal remained constant while the 100 kHz  
148 frequency modulation broadens the spectrum of the radiated signal that was emitted as a high-  
149 frequency electromagnetic field through a horn antenna (Model 3115, ETS Lindgren, Cedar  
150 Park, USA). The antenna was set-up 60 cm above the plants, with an incidence angle of  $45^\circ$   
151 (Fig. 1) and plants were exposed for 30 min in groups of 12, arranged in a rectangular area of  
152 about 20x30 cm. The resulting EMF amplitudes were measured at the plant level using a tri-  
153 axial probe (EP-601, Narda, Cisano sul Neva, Italy) positioned in the center and at each corner  
154 of the plants panel and was found to range from 99 to 110  $\text{V m}^{-1}$  at the corners and 125  $\text{V m}^{-1}$   
155 in the center, Figure 2A). In these conditions, and given the opening of the horn antenna (24 x  
156 13.5 cm), the far field criterion was 95 cm, thus placing the plants close to this distance (about  
157 85 cm at the center of the plate) in the Fresnel zone (between  $D^2/(2*\text{wavelength})$  and

158  $2 \cdot D^2 / \text{wavelength}$ ) to reach the maximum power value. The spectral characteristics of the  
159 EMF (before amplification) was determined by performing a Fast Fourier Transform of the  
160 temporal recording of the EMF signal and showed a typical bandwidth at -20 dBm of about  
161 30 MHz, from 2.43 to 2.46 GHz (Fig. 2B). Arabidopsis plants are rosette-shaped (the floral  
162 stem being most of the time not yet formed), thus creating a more homogeneous biological  
163 system for the interaction with the electromagnetic waves than it would be possible to achieve  
164 with other plants displaying axis with several different orientations (Kundu et al, 2022). The  
165 impact of EMF on plants in terms of thermal effect was assayed through two separate yet  
166 complementary methods. First, we measured the leaf surface temperature using a Luxtron®  
167 reflectometric probe (LumaSense Technologies, Santa Clara, USA). The recording of the  
168 temperature evolution was started after leaf temperature stabilized and performed every 5 s  
169 all along the EMF exposure. Second, we monitor the leaf temperature evolution with a thermal  
170 imaging camera (FLIR E40, Teledyne Technologies, Thousand Oaks, USA) with a built-in  
171 calibration of temperature scale. The Specific Absorption Rate (SAR, in  $\text{WKg}^{-1}$ ) corresponding  
172 to the exposure conditions was determined according to equation (1) :

$$173 \quad \text{SAR} = C \times (dT/dt)_{t \rightarrow 0} \quad (1)$$

174 Where  $C$  is the heat capacity ( $\text{J}^{-1}\text{Kg}^{-1}\text{K}^{-1}$ ),  $T$  the sample temperature and  $t$  the time elapsed  
175 since the beginning of the EMF exposure.

176 Plant leaves were harvested 15, 30 and 60 min after the end of the 30 min exposure,  
177 immediately frozen in liquid nitrogen and lately used for biochemical assays and molecular  
178 biology experiments. Control (*i.e.* non exposed plants) were harvested at the same points of  
179 kinetics (they were manipulated similarly, except that they were not subjected to EMF  
180 exposure). Each sample per condition (control and exposed) and per time point was  
181 constituted with a pool of fully developed leaves from three plants. The sampling of the material  
182 was repeated three times after three independent exposure experiments.

183 *RNA isolation, cDNA synthesis and gene expression quantification*

184 Frozen materials were ground to a fine powder using Retsch® grinder machine prior RNA  
185 isolation using the Nucleospin RNA plant Mini Kit including DNase treatment (Macherey Nagel,  
186 Hoerd, France), according to the manufacturer instructions. The conversion of mRNAs to  
187 cDNA was achieved from 500 ng of total RNA, using the iScript Reverse transcription supermix  
188 for RT-qPCR (Bio-Rad, Marnes-la-Coquette, France) following the manufacturer instruction.  
189 Relative gene expressions were measured using real time quantitative PCR (RT-qPCR)  
190 performed in a final volume of 15 µl of PCR mixture containing 3 µl of 50 times diluted cDNA,  
191 1 µl of primers pairs (10 µM), 4 µl iQ SYBR Green supermix (Bio-Rad, Marnes-la-Coquette,  
192 France) and 7 µl of ultrapure water. The amplifications were performed using the CFX96 Real  
193 Time System (Bio-Rad) with a standard amplification program previously describe in Porcher  
194 et al. (2021). Relative transcript abundances were expressed to the control condition  
195 (unexposed samples harvested just after the end of the exposure) after  $2^{-\Delta\Delta C_t}$  calculation (Livak  
196 and Schmittgen, 2001) along with two house-keeping genes (Aldehyde dehydrogenase, *ALDH*  
197 and Glyceraldehyde-3-phosphate dehydrogenase, *GAPDH*), taken as internal references (Jin  
198 et al., 2019). The primers sequences used in this study are available in the supplementary  
199 table 1.

#### 200 *Hydrogen peroxide quantification*

201 Hydrogen peroxide ( $H_2O_2$ ) levels were assayed using the Amplex™ Red Hydrogen  
202 Peroxide/Peroxidase Assay Kit (Invitrogen). Frozen tissue (20 mg) was homogenized in 100 µl  
203 of 1X kit reaction buffer and kept 5 min on ice. After centrifugation (5 min at 14,000 x g at 4°C),  
204 the supernatant was diluted (10-fold) in 1X reaction buffer. An equal volume of reaction mix  
205 freshly prepared according to the manufacturer protocol and mixed to 50 µl of diluted  
206 supernatant in a flat bottom black plate (Thermo Scientific™ Nunc™ F96 black MicroWell  
207 plate).  $H_2O_2$  levels were determined after a 30 min incubation step (in darkness at room  
208 temperature) by fluorescence measurement (570 nm excitation, 590 nm emission) using a  
209 microplate reader (BMG Labtech FLUOstar Omega) and a standard curve obtained from  
210 known concentrations of commercial  $H_2O_2$  (Sigma).



211 *Quantification of reduced and oxidized forms of glutathione (GSH/GSSG) and ascorbic acid*  
212 *quantifications (AsA/DHA)*

213 Amounts of the reduced (GSH) and oxidized (GSSG) forms of glutathione were determined  
214 from 50 mg of frozen leaf tissues using the GSH-GSSG-Glo™ Assay kit (Promega, Madison,  
215 WI, USA) adapted to plant tissues (Porcher et al., 2020). Ascorbic acid (AsA) and  
216 dehydroascorbic acid (DHA) quantities were determined from 40 mg of frozen tissues using a  
217 colorimetric assay described in Gillespie and Ainsworth (2007).

218 *Lipid peroxidation assay*

219 Malondialdehyde (MDA) content was assayed as described by Jin et al (2020). Briefly, 50 mg  
220 of finely grinded plant tissue was homogenized in 10% trichloroacetic acid and centrifuged for  
221 15 min at 4000xg. The supernatant (300 µl) was added to an equal volume of  
222 0.67% thiobarbituric acid and incubated 30 min at 95°C under gentle shaking in a dry bath.  
223 After a centrifugation (15 min at 10,000xg), the supernatant was used to measure absorbances  
224 at 450, 532 and 600 nm in a BMG Labtech SpectroStar nano spectrophotometer to determine  
225 MDA content (subsequently expressed as µmol.g<sup>-1</sup> FW) accordingly to equation 2:

$$226 \quad [\text{MDA}] = 6.45 \times (A_{532} - A_{600}) - 0.56 \times A_{450} \quad (2)$$

227

228 *Statistical analysis*

229 Statistical analysis was performed with the R statistical software (R core Team, 2020) using  
230 the statistical tests described in figure captions. Graphics were produced using Veusz 3.4  
231 (<https://veusz.github.io/>) software.

## 232 Results

233

### 234 *Leaves temperature evolution after HF-EMF exposure and SAR analysis*

235 The 30 min exposure to the 100 kHz frequency modulated 2.45 GHz HF-EMF did not cause  
236 any significant increase in leaf temperature, as demonstrated by measurements performed  
237 with the Luxtron reflectometric probe (Fig. 3A). The leaves temperature (about 20.3°C)  
238 displayed variations of about 0.1-0.2°C all along the exposure (30 min) and did not show any  
239 significant increase after the application of the EMF exposure (Fig. 3 A, vertical line). Similarly,  
240 image acquisition with a thermal imaging camera showed that the surface temperature of the  
241 leaves was close to 20°C before and after exposure to 2.45 GHz EMF, confirming the  
242 temperature measurement performed with the Luxtron reflectometric probe and the absence  
243 of detectable temperature increase after exposure (Figure 3 B). However, it cannot be  
244 excluded that a slight temperature rise occurred but remained within the temperature  
245 measurement uncertainty (0.1°C). Consequently, we considered a potential maximal leaf  
246 temperature increase of 0.1°C after 30 min of exposure to calculate the corresponding  
247 maximum potential SAR value of 0.21 W kg<sup>-1</sup> according to Eq. (1), assuming a leaf thermal  
248 capacity equal to 3.75 kJ K<sup>-1</sup>kg<sup>-1</sup> (Kitaya *et al.*, 2003).

249

### 250 *Gene expression after plant exposure to EMF*

251 *TCH1* calmodulin transcript remained fairly stable all along the 60 min kinetic in non-exposed  
252 plants (Figure 4A, white bars). In contrast, plant exposure to a non-thermic 100 kHz-modulated  
253 2.45 GHz EMF caused a rapid (15 min) increase in the accumulation of calmodulin transcripts  
254 that became significant 60 min after the end of the exposure (Figure 4A, grey bars). At the  
255 same time, the *ZAT12* transcription factor transcript also rapidly accumulated 15 min after the  
256 end of EMF and became significantly different from that measured in control, unexposed plants  
257 30 and 60 min after the end of the EMF exposure (Figure 4B, grey).

258 We measured the expression level of the NADPH oxidase/respiratory burst oxidase homolog  
259 (RBOH), isoforms D and F, that are key enzymes involved in the production of superoxide  
260 rapidly dismutates into  $H_2O_2$  by apoplastic superoxide dismutases (SODs), in response to  
261 environment constraints (Castro et al., 2021). The expression level of *RBOHD* gene isoform  
262 remained essentially constant and unaffected after EMF exposure (Figure 5A). In contrast, if  
263 the accumulation of the *RBOHF* isoform transcript displayed slight changes in the control,  
264 unexposed plants (Figure 5B, white bars), the exposed plants showed constantly higher  
265 amounts of transcripts that became statistically significant 60 min after the end of the exposure  
266 (Figure 5B, gray bars), suggesting a higher potential of  $H_2O_2$  production in exposed plants.  
267 The expression of the main catalase gene isoform *CAT2* involved in the bulk scavenging of  
268  $H_2O_2$  (Mhamdi et al., 2010), was constant in control plants and not primed by EMF exposure  
269 (Figure 5C): a significant decrease in the accumulation of *CAT2* transcript was even noted  
270 30 min after the end of the exposure. In contrast, the expression level of the ascorbate  
271 peroxidase *APX1*, involved in the fine tuning of  $H_2O_2$  homeostasis was significantly higher in  
272 exposed plants after 60 min (Figure 5D, grey bars). It is worth noting that it remained  
273 essentially constant in control, unexposed plants (Figure 5D, white bars).

274

#### 275 *H<sub>2</sub>O<sub>2</sub> accumulation and scavenging*

276 Hydrogen peroxide significantly accumulated 60 min after the end of the exposure,  
277 comparatively to that observed in the unexposed control (Figure 6A). The amount of  $H_2O_2$  was  
278 about  $213.8 \text{ nmol.g}^{-1} \text{ FW}$  while it increased to  $293.9 \text{ nmol.g}^{-1} \text{ FW}$  (+37.4 %) in the exposed  
279 plants.

280 The amount of total glutathione increase (however not significantly) in exposed plants 60 min  
281 after the exposure (Figure 6B, left), mainly due to an elevation in the content of GSH (about  
282 40%, from  $174.52$  to  $245.53 \text{ nmol g}^{-1} \text{ FW}$  in control and exposed plants, respectively), while  
283 the amount of GSSG remains approximately the same ( $21.47$  vs  $26.11 \text{ nmol g}^{-1} \text{ FW}$  in control

284 and exposed plants, respectively). The amplitude of these changes was not sufficient to  
285 significantly shift the GSH/GSSH ratio (8.64 vs 9.31 in control and exposed plants,  
286 respectively, Fig 6B right). The quantification ascorbic acid (ASA) slightly increased (about  
287 19.6%, from 11.34 to 13.57  $\mu\text{mol g}^{-1}$  FW) after exposure, while being not significantly different  
288 from that found in unexposed plants (Fig 6C, right). In contrast, the quantity of dehydroascorbic  
289 acid (DHA) was significantly higher (about 124%, from 1.95 to 4.38  $\mu\text{mol g}^{-1}$  FW) 60 min after  
290 exposure (Fig 6C, right).

291

### 292 *Malondialdehyde formation*

293 The amount of malondialdehyde (Fig 6D) remained very similar in control and exposed plants  
294 (4.88 vs 4.67  $\mu\text{mol. g}^{-1}$  FW, respectively) showing that exposing plants for 30 min to a non-  
295 thermic 100 kHz-modulated 2.45 GHz EMF did not affect lipid peroxidation.

296

### 297 **Discussion**

298 The existence of cellular/molecular and morphological changes after non-thermal  
299 electromagnetic exposure is largely a matter of debate in the present literature. An EMF-  
300 exposure is generally considered as non-thermal if the tissue temperature rise resulting from  
301 the exposure is lower than 1°C after 30 min of exposure (IEEE, 2005; ICNIRP, 2020). Although  
302 human cells did not show any significant metabolic changes after such kind of exposure (Roux  
303 et al., 2010; Habauzit et al., 2014), the situation is far from being so straightforward in plants.  
304 Indeed, several works highlighted that electromagnetic exposures of various frequencies and  
305 consistent SAR levels (about 0.5  $\text{Wkg}^{-1}$ ) caused different kind of cellular responses, including  
306 cellular changes and mitotic alteration (Vian et al., 2016; Kaur et al., 2021). Răcuciu et al.  
307 (2015) showed that exposing maize seedlings to 1 GHz for 8 h (DAS of 0.47  $\text{Wkg}^{-1}$ ) resulted  
308 in an important decrease in the total pigment (a + b chlorophylls and carotenoids) and a  
309 reduction of plantlets growth. Akbal et al (2012) demonstrated that 48 h exposure of dormant  
310 *Lens culinaris* seeds to HF-EMF (1800 MHz, SAR 0.76  $\text{W kg}^{-1}$ ) did not affect germination but

311 caused a delayed root growth inhibition and an increase in c-mitosis number. Other DNA  
312 alterations (increase in micronuclei, ranging from 2.3 to 7-fold compared to non-exposed  
313 samples) were found after exposing *Vicia faba* root tips to 915 MHz (SAR: 0.4-1.6 W kg<sup>-1</sup>)  
314 during 72 h (Gustavino et al., 2016). Pesnya and Romanovsky (2013) also observed an  
315 increase in mitotic abnormalities after exposing onion root tips to 900 MHz for 9 h (SAR:  
316 1.4 W kg<sup>-1</sup>). Although it is likely that such alterations are accompanied by changes in gene  
317 expression, relatively few studies have investigated this aspect. Engelmann et al. (2008)  
318 analyze the global gene expression pattern based on Affymetrix ATH1 microarrays after  
319 exposing cultured *Arabidopsis thaliana* cells for 24h to HF-EMF (1.9 GHz, average SAR:  
320 0.75 Wkg<sup>-1</sup>) and found that 10 genes displayed a significant yet limited variation in their  
321 expression (2.5 fold).

322 Here we used a horn antenna (Fig. 1) to generate an electromagnetic field whose  
323 characteristics were deterministic with a fixed incidence, contrary to a finite number of plane  
324 waves with random polarization and incidence angle, as in the MSRC that we used in our  
325 former studies (Beaubois et al., 2007; Grémiaux et al., 2016; Roux et al., 2008; Vian et al.,  
326 2006). This point was essential to ensure that all the waves contributed to the coupling with  
327 the plants (here represented by the rosettes of *Arabidopsis* on which the waves reached with  
328 an angle of incidence of 45°, Fig. 1). We used in the present work an exposure condition that  
329 keep constant the amplitude of the electromagnetic wave (2.45 GHz, about 100 Vm<sup>-1</sup> at the  
330 plant level, Fig. 2A) while the 100 kHz frequency modulation broadens the spectrum of the  
331 radiated signal (about 30 MHz bandwidth, Fig. 2B). This exposure condition did not cause any  
332 measurable shift in leaf surface temperature over the 30 min of the exposure duration (the  
333 temperature oscillation was less than 0.2 °C), as reported by two different measurement  
334 methods: the Luxtron reflectometric probe and the thermal imaging (Fig. 3A-B). This absence  
335 of tissue heating, that allowed us to selectively study the non-thermal responses of plants to  
336 HF-EMF exposure, may be related to the high surface to volume ratio of the plants (Vian et al.,  
337 2007) that strongly favor heat dissipation. One could note that the resolution of the thermal

338 camera sensor does not exclude the existence of microscopic temperature rise that are  
339 however unlikely to affect the whole leaf temperature.

340 Taking the uncertainty of the temperature measurement as the maximum potential heating  
341 value (here 0.1°C) resulted in a maximal potential SAR value of 0.21 W kg<sup>-1</sup>, assuming a leaf  
342 thermal capacity equal to 3.75 kJ K<sup>-1</sup>kg<sup>-1</sup> (Kitaya *et al.*, 2003). It is worth noting that this  
343 potential maximal value was also comparable to those measured in other works reporting  
344 biological effects after exposing plants to HF-EMF: 0.169 Wkg<sup>-1</sup> (Kumar *et al.*, 2016), 0.282 W  
345 kg<sup>-1</sup> (Chandel *et al.*, 2017), for which no temperature increase was neither observed.

346 We reported here, in this non-thermal exposure condition, that the expression of stress-related  
347 genes such as calmodulin *TCH1* (Fig. 4A) or the transcription factor *ZAT12* (Fig. 4B)  
348 significantly increased 60 min after the end of the exposure, while remained at a low level in  
349 the control, unexposed plants. *TCH1* gene expression was previously reported to increase in  
350 tomato after exposure to non-thermal HF-EMF (Roux *et al.*, 2006; Roux *et al.*, 2008) (,a result  
351 later confirmed by Rammal *et al.* (2014) also in tomato and more recently by Kundu *et al.*  
352 (2021a,b) in rice. The expression level of stress-related transcription factor gene *bZIP* was also  
353 reported to increase in such condition (Vian *et al.*, 2006; Beaubois *et al.*, 2007), a result also  
354 recently confirmed after exposing rice to low power HF-EMF (Kundu *et al.*, 2021a,b). It is worth  
355 noting that the accumulations of *TCH1*, *bZIP* and *PIN2* occurred more rapidly (0-15 min after  
356 the end of the 10 min exposure) when the exposure was performed in a MSRC (Roux *et al.*,  
357 2006, 2008; Beaubois *et al.*, 2007). This difference may result from the multiple reflections of  
358 the electromagnetic waves on the metallic walls of the CRBM, which considerably increase the  
359 possibilities of coupling of the electromagnetic field with the plant. Changes in the expression  
360 levels of calmodulin (*TCH1*) and *ZAT12* genes suggested an activation of calcium and ROS  
361 signal transduction pathways that are closely related (Davletova *et al.*, 2005; Gilroy *et al.*,  
362 2016). We therefore analyzed the expression level of several genes corresponding to the main  
363 actors of production and scavenging of H<sub>2</sub>O<sub>2</sub>. We showed that the expression level of the  
364 *RBOHD* gene, involved in hypoxia signaling (Yang and Hong, 2015) was not affected after

365 exposing plants to HF-EMF (Fig. 5A). In contrast, *RBOHF*, which appears as a regulation hub  
366 targeted by calcium-inducible protein kinases (Han et al, 2018; Castro et al., 2021), was  
367 induced after exposure and became significantly more expressed after 60 min (Fig. 5B),  
368 suggesting an increase in H<sub>2</sub>O<sub>2</sub> production and confirmed by its direct quantification (Fig. 6A).  
369 Interestingly, this was concomitant with the increased expression of *TCH1* (Fig. 4A) that is a  
370 marker of an increase in cytosolic calcium concentration.

371 The expression level of catalase (*CAT2*), an essential pathway of bulk, low affinity H<sub>2</sub>O<sub>2</sub>  
372 detoxification (Gill and Tuteja, 2010), was not affected by HF-EMF exposure (Fig. 5C),  
373 whereas that of *APX1*, which finely regulates the H<sub>2</sub>O<sub>2</sub> level due to its high affinity, was  
374 significantly increased after 60 min (Fig. 5D), indicating a possible involvement of the  
375 AsA/glutathione detoxification pathway. The total quantity of glutathione did not change after  
376 exposure (Fig. 6B), as well as the GSH/GSSH ratio. Ascorbic acid (Fig. 6C, left) level did not  
377 change either, while the dehydroascorbic acid was the only actor of the AsA/glutathione  
378 pathway displaying a significant variation 60 min after exposure (Fig. 6C, right). Taken  
379 together, these results showed that non-thermic HF-EMF trigger a low but significant increase  
380 in H<sub>2</sub>O<sub>2</sub> production which reflects more of a signal transduction mechanism, acting in synergy  
381 with possible calcium movements as reported by calmodulin increase, than a stress response.  
382 This observation is consistent with the general scheme proposed by Kaur et al. (2021) in which  
383 the initial interaction of electromagnetic field with plants triggers both calcium movements and  
384 ROS metabolism as initial events of signal transduction cascade that later cause changes in  
385 plant development. Modifications in ROS production and/or scavenging were in fact reported  
386 among others in wheat (Chen et al., 2009), mung bean (Singh et al., 2012), onions (Chandel  
387 et al., 2017), duckweed (Tkalec et al., 2007) and *Arabidopsis* (Senevirathna et al., 2020). It is  
388 also striking that quite similar changes in ROS metabolism have been reported after exposure  
389 of plants to a static magnetic field (SMF) or to an extremely low frequency electromagnetic  
390 field (ELF-EMF), despite the fundamental differences in the nature of these fields. Indeed,  
391 exposing plants to static magnetic field higher than the geomagnetic field generally caused an



392 increase in H<sub>2</sub>O<sub>2</sub> production and ROS-scavenging enzymes activities and H<sub>2</sub>O<sub>2</sub> synthesis-  
393 related gene expression (see the reviews from Maffei et al., 2014, Saletnik et al., 2022 and  
394 Hafeez et al. (2023) for details). In contrast, exposing plants to near null magnetic field (much  
395 lower to that of the geomagnetic field) resulted in a drastic reduction of H<sub>2</sub>O<sub>2</sub> in roots and shoots  
396 (5-folds after 24h), likely related to the up-regulation of gene expression of ROS scavenging  
397 enzymes (Parmagnani et al., 2022). ROS metabolism is therefore an essential component in  
398 plants' perception of the presence/absence of static magnetic fields, as well as of their  
399 exposure to very high frequency electromagnetic fields.

## 400 **Conclusion**

401 We demonstrated here that a very high frequency electromagnetic field, which we have verified  
402 as having no heating effects in plants, was able to induce rapid changes in the expression of  
403 several genes involved in signal transduction or ROS metabolism, as well as the production of  
404 hydrogen peroxide. This ability appears to be unique to plants, as it is clear that non-thermal  
405 exposure does not elicit a molecular response in humans (Roux et al., 2010; Habauzit et al,  
406 2014). These results, in addition to many previous observations, raise the question of the  
407 relevance of SAR in plants, insofar as this quantity (which otherwise adequately accounts for  
408 biological effects in animals) is based on the thermal effects induced by exposure to  
409 electromagnetic fields. This major difference could be the consequence of the fundamental  
410 contrast in the logic of development (Vian et al., 2007), as plants create large surfaces, which  
411 forces them to immobility but favors interaction with the environment (in this case interaction  
412 with electromagnetic waves), whereas animals create volume and comparatively little surface  
413 with the environment (which favors their mobility). The interaction mechanism of the  
414 electromagnetic field, which clearly does not depend on thermal effects, remains as yet  
415 misunderstood although several hypotheses have been proposed (Cifra et al, 2011;  
416 Romanenko et al, 2017). Studies are still needed, and we believe that a more comprehensive  
417 knowledge of the molecular and biochemical changes (such as those provided by an RNA  
418 sequencing or metabolic analyses) taking place after non-thermal exposure could allow us to



419 make effective progress in understanding the mechanism of interaction between plants and  
420 electromagnetic field.

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422

423 **Figure legends**

424

425 **Fig. 1:** experimental set-up. The electric polarization is indicated by a red arrow.

426 **A.** Schematic representation. The electromagnetic field was generated by a signal synthesizer,  
427 amplified, and then emitted through a horn antenna placed at a distance  $h=60$  cm above the  
428 plants with an angle of  $45^\circ$ .

429 **B.** Picture of the set-up. Note that the experiments are performed in a shielded room, totally  
430 isolated from the electromagnetic background noise.

431

432 **Fig. 2:** electromagnetic field characteristics

433 **A.** Field amplitude in the plant panel. The field amplitude was measured using a Narda EP-  
434 601 triaxial probe in the center and at each corner of the plant panel.

435 **B. Frequency** Bandwidth of the exposure. A Fast Fourier Transform of the temporal signal  
436 revealed that the bandwidth of about 30 MHz of the signal is centered on the fundamental  
437 frequency (2.45 GHz) .

438

439 **Fig. 3:** Evolution of plant leaf temperature after EMF exposure.

440 **A.** The leaf surface temperature was continuously recorded from 2 min before the  
441 application of the 100 kHz-modulated 2.45 GHz EMF field (a), during the 30 min  
442 exposure between the two red vertical lines (b) and 3.5 min after the exposure stop (c).  
443 The measurement was performed using a Lumasense Luxtron modèle STS  
444 reflectometric probe. Representative recording out of 3 independent experiments.

445 **B.** Thermic imaging of the plant's panel before (left) of after (right) the application of a  
446 30 min 100 kHz-modulated 2.45 GHz EMF. The red cross materializes the possibility

447 to measure selectively the temperature on a very reduced surface of the leaf.  
448 Positioned on hot spots present on both unexposed and exposed samples, the  
449 measures revealed similar surface temperatures (24.3°C and 23°C, on unexposed and  
450 exposed samples, respectively).

451

452 **Fig. 4:** Effect of HF-EMF exposure on signal transduction-related gene expression

453 Relative transcript accumulations of *TCH1* calmodulin (**A**) and *ZAT12* transcription factor (**B**)  
454 genes, both implicated in rapid responses of plants to various kind of environmental stimuli, in  
455 unexposed (control) and exposed plants (15, 30 and 60 min after 30 min exposure to HF-EMF  
456 at 2.45 GHz, SAR<0.21 WKg<sup>-1</sup>). Data are means of n = 3 biological independent replicates ±  
457 s.e. Asterisks indicate significant differences after the non-parametric Wilcoxon-Mann-Whitney  
458 test at the 95% confidence interval (\*: P < 0.05; \*\*: P < 0.01 and \*\*\*: P < 0.001) between the  
459 exposed and control condition for each different time point.

460

461 **Fig. 5:** effect of HF-EMF exposure on ROS-related gene expression

462 Relative transcript accumulation of (**A**) NADPH oxidases *RBOHD* and (**B**) *RBOHF*,  
463 (**C**) catalase *CAT2* and (**D**) ascorbate peroxidase *APX1* genes, all implicated in H<sub>2</sub>O<sub>2</sub>  
464 production and scavenging, in unexposed (control) and exposed plants (15, 30 and 60 min  
465 after 30 min exposure to HF-EMF at 2.45 GHz, SAR<0.21 WKg<sup>-1</sup>). Data are means of n = 3  
466 biological independent replicates ± s.e. Asterisks indicate significant differences after the non-  
467 parametric Wilcoxon-Mann-Whitney test at the 95% confidence interval (\*: P < 0.05; \*\*: P <  
468 0.01 and \*\*\*: P < 0.001) between the exposed and control condition for each different time  
469 point.

470

471

472 **Fig. 6:** effect of HF-EMF exposure on ROS metabolism and MDA production

473 Amounts of **(A)** H<sub>2</sub>O<sub>2</sub>, **(B)** total glutathione (GSH, GSSG) and GSH/GSSG ratio, **(C)** amounts  
474 of ascorbic (ASA) and dehydroascorbic (DHA) acids and **(D)** amounts of malondialdehyde  
475 (MDA) in *Arabidopsis thaliana* leaves in unexposed (control) and exposed plants (60 min after  
476 30 min exposure to HF-EMF at 2.45 GHz, SAR<0.21 WKg<sup>-1</sup>). Data are means of n = 3  
477 biological independent replicates ± s.e. Asterisks indicate significant differences after the non-  
478 parametric Wilcoxon-Mann-Whitney test at the 95% confidence interval (\*: P < 0.05; \*\*: P <  
479 0.01 and \*\*\*: P < 0.001) between the exposed and control condition.

480

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484 **Table S1: List of primer sequences used in this study.**

485 Genes in bold were used as housekeeping genes

Gene ontology	Gene name	Locus id	Primer sequences	Amplicon size	References
<b>Glyceraldehyde-3-phosphate dehydrogenase</b>	<i>GAPDH</i>	At1G13440	Forward: TCAGGAACCCTGAGGACATC Reverse: CGTTGACACCAACAACGAAC	174 bp	Jin <i>et al.</i> (2019)
<b>Aldehyde dehydrogenase</b>	<i>ALDH</i>	At3G66658	Forward: TGGGATATTTCCCTGCTCTG Reverse: GAAGAATCCGCAAGAACTGC	123 bp	Jin <i>et al.</i> (2019)
Calmodulin 1	<i>TCH1</i>	At5G37780	Forward: ATGGAAACGGCACTATCGAC Reverse: ATTGGTCATCACATGGCGTA	161 bp	newly designed
Zinc finger protein/stress responses	<i>ZAT12</i>	At5G59820	Forward: TGCGAGTCACAAGAAGCCTA Reverse: GTGTCCTCCCAAAGCTTGTC	127 bp	newly designed
Respiratory burst oxidase homolog D	<i>RBOHD</i>	At5G47910	Forward: CGTGGAGATCACGCTAGACA Reverse: AGGAGGTGGTGTGTTGAGG	149 bp	newly designed
Respiratory burst oxidase homolog F	<i>RBOHF</i>	At1G64060	Forward: GGTGCACCAGCACAAGATTA Reverse: TGAAATCCGAGATCGAATCC	148 bp	newly designed
Catalase 2	<i>CAT2</i>	At4G35090	Forward: CGAGGTATGACCAGGTTTCGT Reverse: CTCCAGGCTCCTTGAAGTTG	114 bp	newly designed
Ascorbate peroxidase 1	<i>APX1</i>	At1G07890	Forward: GCACTATTGGACGACCCTGT Reverse: AGCAAACCCAAGCTCAGAAA	117 bp	newly designed

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492

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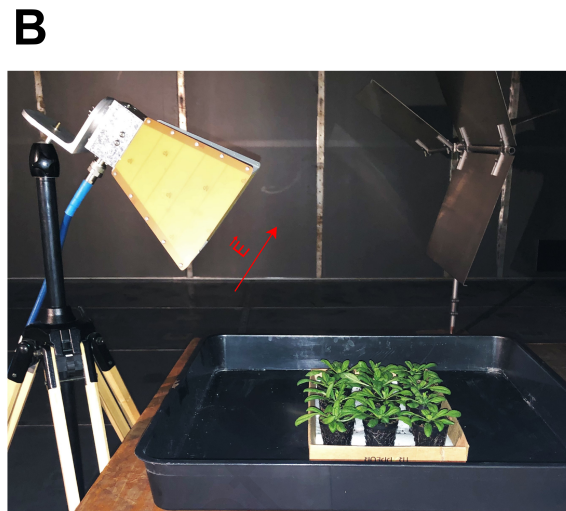
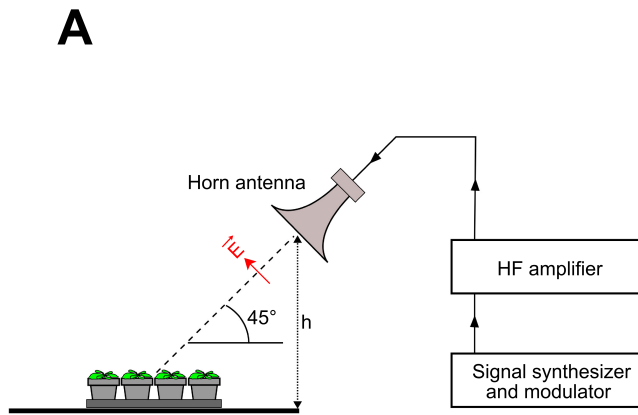
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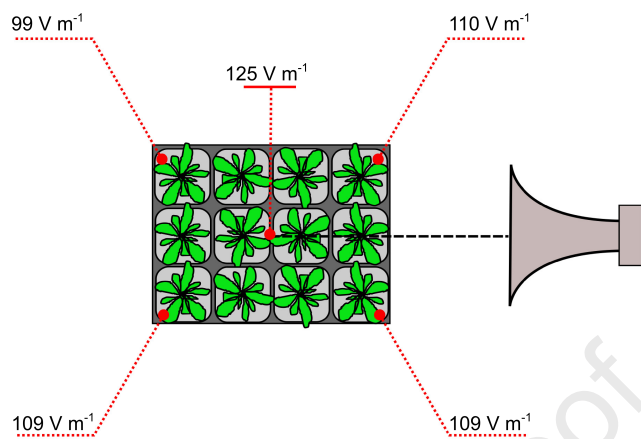
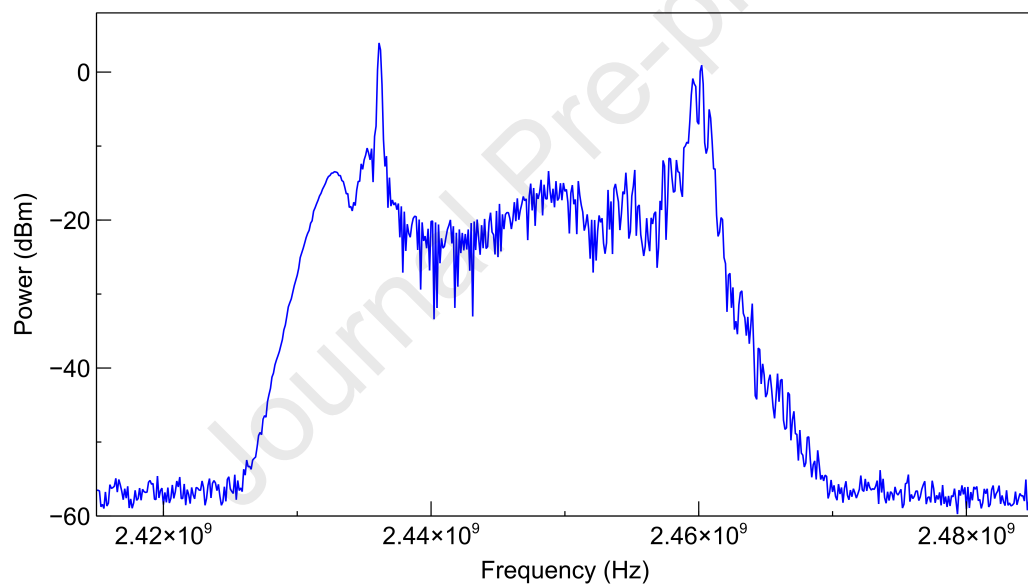
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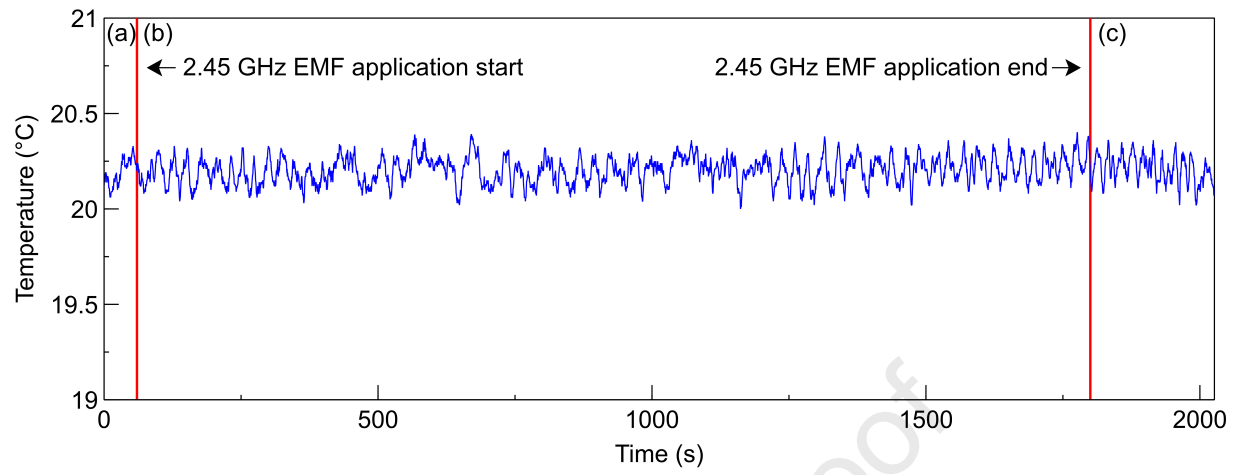
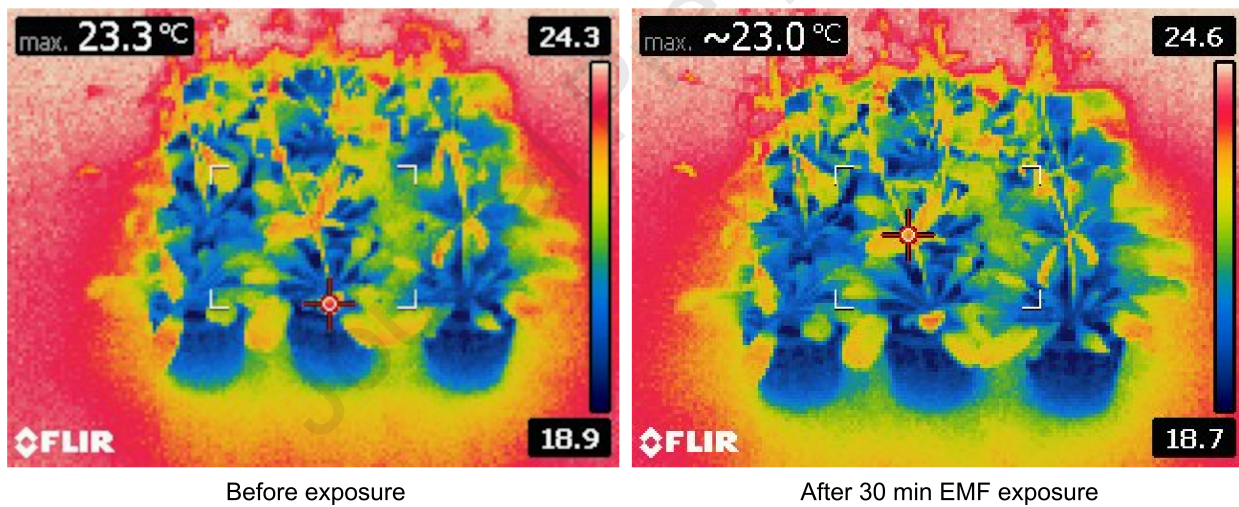
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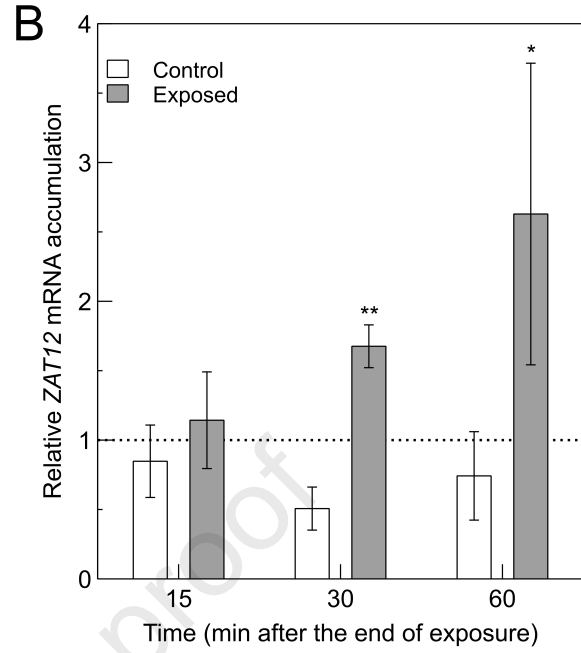
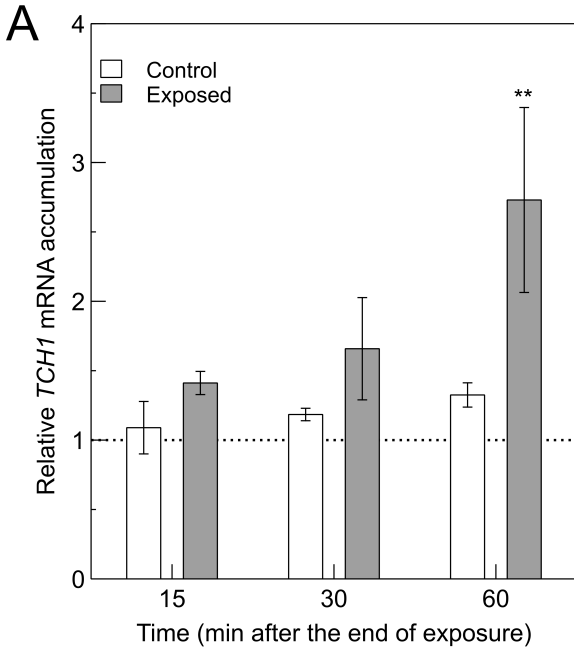
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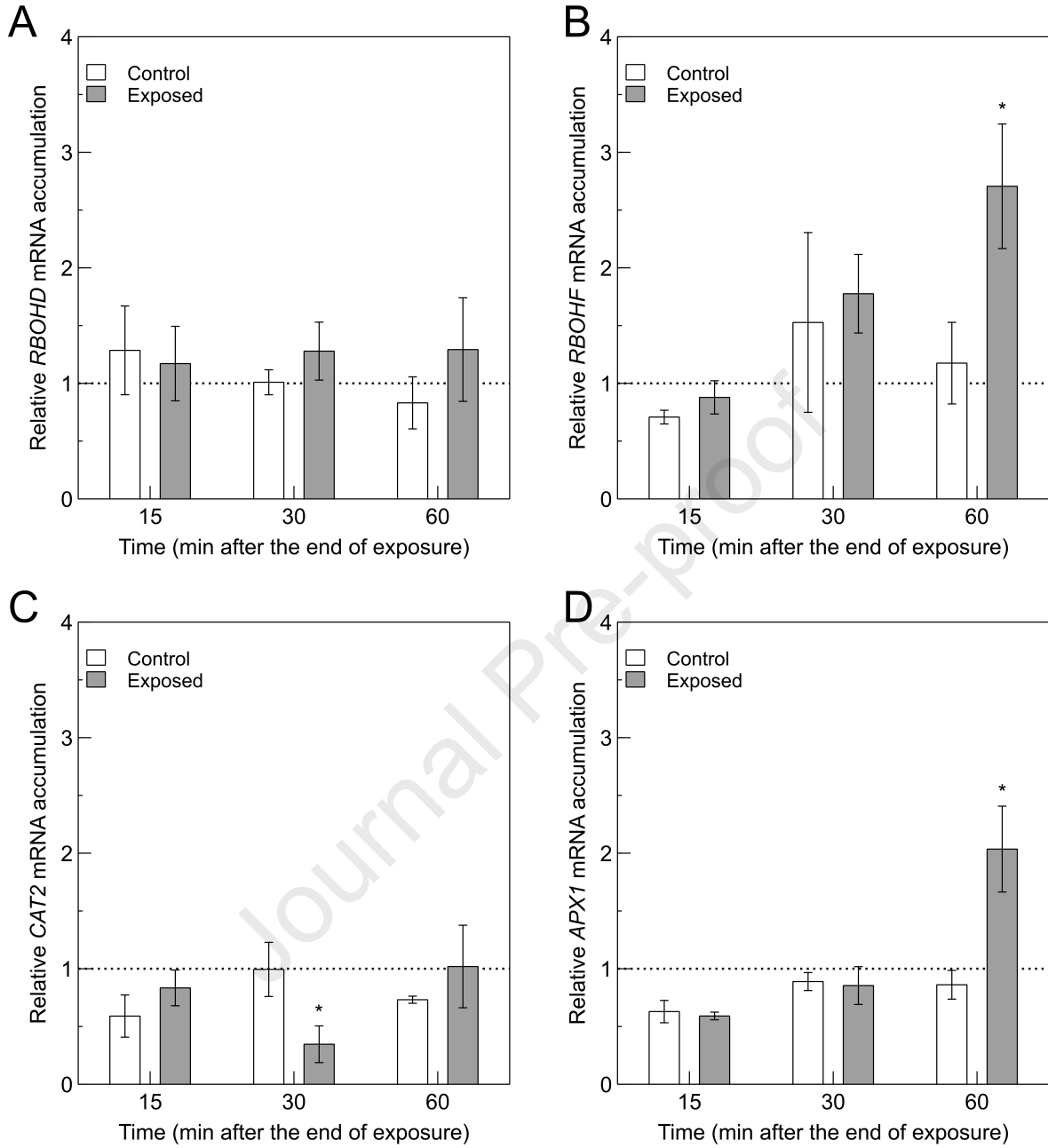


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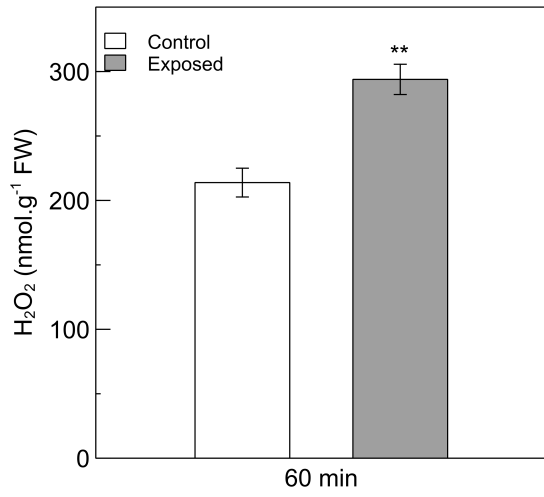
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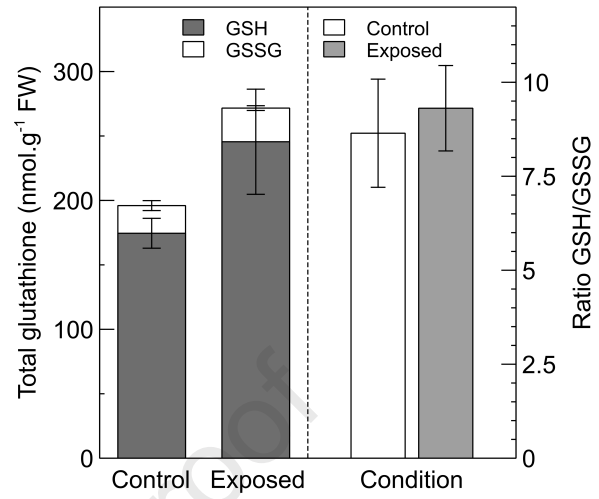




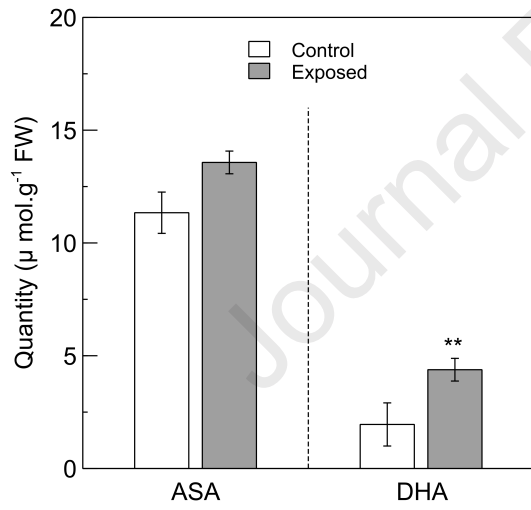
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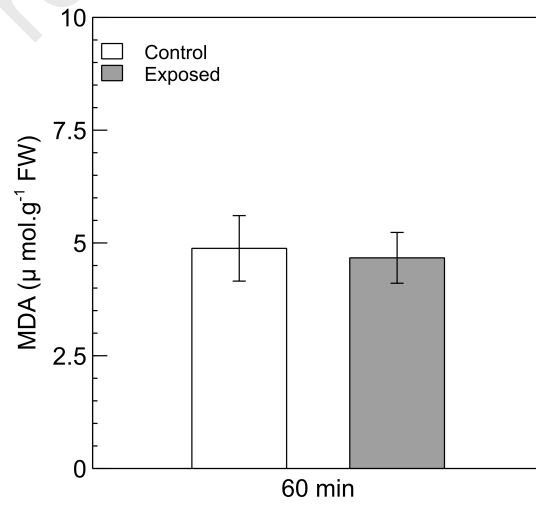
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## Highlights

- In our condition, 2.45 GHz HF-EMF exposure did not induce thermal effect in plants
- We observed rapid and transient changes of gene expression and H<sub>2</sub>O<sub>2</sub> metabolism
- Non-thermal biological responses to EMF exposure do exist in plants

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**Declaration of interests**

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests:

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