

# Inheritance of heat tolerance in perennial ryegrass (*Lolium perenne*, Poaceae): evidence from progeny array analyses

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**Background:** Heat stress is considered one of the most important environmental factors influencing plant physiology, growth, development, and reproductive output. The occurrence and damage caused by heat stress will likely increase with global climate change. Thus, there is an urgent need to better understand the genetic basis of heat tolerance, especially cool season plants.

**Materials and methods:** In this study, we assessed the inheritance of heat tolerance in perennial ryegrass (*Lolium perenne* L. subspecies *perenne*), a cool season grass, through a comparison of two parental cultivars with their offspring. We crossed a heat tolerant cultivar (Kangaroo Valley) with a heat sensitive cultivar (Norlea), to generate 72 F1 progeny. Both parents and their progeny were then exposed to heat stress for 40 days, and their photosynthetic performance (Fv/Fm values) and leaf H<sub>2</sub>O<sub>2</sub> concentrations were measured.

**Results:** As expected, Kangaroo Valley had significantly higher Fv/Fm values and significantly lower H<sub>2</sub>O<sub>2</sub> concentrations than Norlea. For the F1 progeny, values of Fv/Fm decreased gradually with increasing exposure to heat stress, while the content of H<sub>2</sub>O<sub>2</sub> increased. The progeny had a wide distribution of Fv/Fm and H<sub>2</sub>O<sub>2</sub> values at 40 days of heat stress. Approximately 90% of the 72 progeny had Fv/Fm values that were intermediate to the values of the two parental cultivars and 67% of the progeny had H<sub>2</sub>O<sub>2</sub> concentrations intermediate to their two parents.

**Conclusion:** Results of this study indicate considerable additive genetic variation for heat tolerance within the progeny array generated from this cross, and such diversity could be used to improve heat tolerance in perennial ryegrass cultivars. Our findings point to the benefits of combining physiological measurements within a genetic framework to assess the inheritance of heat tolerance, a complex phenotypic trait.

1 **Inheritance of heat tolerance in perennial ryegrass**  
2 **(*Lolium perenne*, Poaceae): evidence from progeny**  
3 **array analyses**

4  
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23

24 **Abstract**

25 Heat stress is considered one of the most important environmental factors influencing plant

26 physiology, growth, development, and reproductive output. The occurrence and damage caused

27 by heat stress will likely increase with global climate change. Thus, there an urgent need to better

28 understand the genetic basis of heat tolerance, especially cool season plants. In this study, we

29 assessed the inheritance of heat tolerance in perennial ryegrass (*Lolium perenne* L. subspecies

30 *perenne*), a cool season grass, through a comparison of two parental cultivars with their

31 offspring. We crossed a heat tolerant cultivar (Kangaroo Valley) with a heat sensitive cultivar

32 (Norlea), to generate 72 F1 progeny. Both parents and their progeny were then exposed to heat

33 stress for 40 days, and their photosynthetic performance (Fv/Fm values) and leaf H<sub>2</sub>O<sub>2</sub>  
34 concentrations were measured. As expected, Kangaroo Valley had significantly higher Fv/Fm  
35 values and significantly lower H<sub>2</sub>O<sub>2</sub> concentrations than Norlea. For the F1 progeny, values of  
36 Fv/Fm decreased gradually with increasing exposure to heat stress, while the content of H<sub>2</sub>O<sub>2</sub>  
37 increased. The progeny had a wide distribution of Fv/Fm and H<sub>2</sub>O<sub>2</sub> values at 40 days of heat  
38 stress. Approximately 90% of the 72 progeny had Fv/Fm values that were intermediate to the  
39 values of the two parental cultivars and 67% of the progeny had H<sub>2</sub>O<sub>2</sub> concentrations  
40 intermediate to their two parents. Results of this study indicate considerable additive genetic  
41 variation for heat tolerance within the progeny array generated from this cross, and such  
42 diversity could be used to improve heat tolerance in perennial ryegrass cultivars. Our findings  
43 point to the benefits of combining physiological measurements within a genetic framework to  
44 assess the inheritance of heat tolerance, a complex phenotypic trait.

45

## 46 **Introduction**

47 Based on the various habitats they occupy, plants require certain environmental conditions to  
48 maintain the abundance and persistence of their populations ([Harper, 1977](#)). During their  
49 lifetime, however, most plants experience abiotic stress when exposed to unfavorable chemical  
50 and physical environmental conditions such as heavy metals, high salinity, excessive solar  
51 radiation, freezing temperatures, severe drought, and extremely high temperatures ([Nilsen and  
52 Orcutt, 1996](#)). Of these stressors, drought and heat stress are among the two most important  
53 environmental factors influencing plant physiology, growth, development, and reproductive  
54 output (i.e., yield) ([Jiang and Huang, 2001](#); [Prasad et al., 2008](#), [Jespersen et al., 2017](#)).

55 According to [Wahid et al. \(2007\)](#), heat stress (or heat shock) in plants occurs when  
56 temperatures rise above a threshold level for sufficient time to result in irreversible damage to  
57 plant growth and development. Although heat stress usually occurs with an increase in  
58 temperature of 10-15 °C above ambient; heat stress is also influenced by the intensity, duration,  
59 and rate of increase in temperature ([Wahid et al., 2012](#)). Thus, heat stress in plants can occur on a  
60 daily or seasonal basis and can vary from year-to-year. In addition, the occurrence and damage  
61 caused by heat stress will likely increase with global climate change ([Walter et al., 2013](#); [Bita  
62 and Gerats, 2013](#)). Due to human activities, substantial increases in the concentration of  
63 greenhouse gases are occurring, and global air temperatures are predicted to increase 1-4.5 °C  
64 above the current level by 2100, depending on different carbon emission scenarios ([Rogelj et al.,  
65 2012](#), [IPCC, 2019](#)). Moreover, human-caused climate change is also associated with extreme  
66 climate events such as precipitation extremes, flooding, frosts, drought, and excessive heat ([Niu  
67 et al., 2014](#); [Stott, 2015](#)). Thus, future climate change is expected to cause serious damage to the  
68 growth and yield of native plants and crop plants, especially C<sub>3</sub> plants and crops ([Lobell and  
69 Field, 2007](#); [Wahid et al., 2012](#)).

70 Exposure of plants to excessively high temperatures can result in a range of complex  
71 responses from molecular and cellular, to whole plant levels ([Baniwal et al., 2004](#); [Kotak et al.,  
72 2007](#); [Wahid et al., 2012](#); [Prasad et al., 2008](#), [Mittler et al., 2011](#); [Soliman et al., 2011](#)). Once leaf  
73 temperatures rise above a threshold level (35 to 40 °C for most plants), protein denaturation and  
74 loss of cell membrane fluidity begins to take place and cell damage and programmed cell death  
75 may occur ([Huang and Xu, 2008](#); [Horvath et al. 2012](#)). Depending on the intensity and duration  
76 of exposure to high temperatures, plant tissue type, and phenological stage, heat stress in plants  
77 can induce the following responses: 1) loss of cell water content, 2) reduced photosynthetic

78 activity, 3) oxidative stress, 4) scorching of tissues and premature leaf senescence and abscission,  
79 5) reduced growth rates through inhibition of shoot and root growth, 6) damage or alteration of  
80 floral (reproductive) tissues, and 7) reduced seed number and quality (see Figure 1 in  
81 [Hasanuzzaman et al., 2013](#)).

82 With heat stress, reductions in photosynthetic activity and efficiency may take place because  
83 high temperatures can lead to the dissociation or inhibition of oxygen evolving complexes (OEC)  
84 and reduce the activity of photosystem II (PSII) ([Wahid et al., 2007](#)). Photosynthetic  
85 performance during heat stress can be quantified by measuring chlorophyll fluorescence  
86 parameters ([Baker and Oxborough 2004](#); [Rosyara et al., 2010](#)). One such parameter is Fv/Fm,  
87 which is calculated as the ratio between variable fluorescence ( $F_v = F_m - F_o$ ) and maximum  
88 fluorescence (Fm). Exposure to high temperatures can also induce oxidative stress in plants by  
89 uncoupling enzymes and metabolic pathways which generates reactive oxygen species (ROS)  
90 that can damage multiple cellular organelles and physiological processes ([Locato et al. 2008](#);  
91 [Soliman et al., 2011](#); [2012](#); [Hasanuzzaman et al., 2013](#)). Reactive oxygen species (ROS) include  
92 singlet oxygen, superoxide radical, hydroxyl radical, and hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>). Hydrogen  
93 peroxide is the most stable ROS and its adverse effects in plants include membrane lipid  
94 peroxidation, toxicity, and cell death. Interestingly, recent studies have highlighted the important  
95 role of H<sub>2</sub>O<sub>2</sub> as a signaling molecule in plants, triggering tolerance responses to abiotic stresses  
96 ([Suzuki et al., 2012](#); [Baxter et al., 2014](#)).

97 Plants are sessile organisms that cannot evade abiotic stressors. Thus, they have developed  
98 several mechanisms for mitigating and surviving heat stress (see Figure 4 in [Hasanuzzaman et](#)  
99 [al., 2013](#)). These mechanisms include short-term avoidance or acclimation mechanisms and  
100 long-term phenological and morphological adaptive traits such as early maturation, enhanced

101 root density and depth, changing leaf orientation and leaf rolling, transpirational cooling, and/or  
102 alteration of membrane lipid compositions (Wahid et al., 2007; 2012; Prasad et al., 2008;  
103 Hasanuzzaman et al., 2013; Jespersen et al., 2017). Additionally, plants have developed  
104 molecular, cellular, and physiological adaptations for tolerating heat stress (Wahid et al., 2012;  
105 Hasanuzzaman et al., 2013). These include signaling cascades and regulation of gene expression  
106 by transcription factors (Yang et al., 2014; Ohama et al., 2016; Jespersen et al., 2017), expression  
107 of heat shock proteins (HSPs) and molecular chaperones (Horvath et al., 2012; Davies et al.,  
108 2018), enzymatic and non-enzymatic antioxidant defense to prevent the harmful effects of ROS  
109 (Gulen and Eris, 2004), and the production of osmo-protectants or compatible solutes (Wahid et  
110 al., 2007; Hasanuzzaman et al., 2013). Clearly, heat tolerance in plants is controlled by a  
111 complex set of many genes and interacting mechanisms, and not just a single gene or mechanism  
112 (Erdayani et al., 2020).

113 Perennial ryegrass (*Lolium perenne* L. subspecies *perenne*) is a cool season ( $C_3$ ), perennial  
114 grass that has a caespitose (bunch) growth form and can grow to a height of approximately 10-90  
115 cm. Perennial ryegrass originated in the Middle East, and then dispersed across Europe and  
116 North Africa with the spread of agriculture (Balfourier et al. 2000). Perennial ryegrass has  
117 subsequently been introduced around the globe and is considered a weed, or an invasive species,  
118 in natural communities in many regions. It is also one of the most common pasture grasses in  
119 temperate climates regions where it is used as a forage grass for livestock and for hay  
120 production. In addition, it is widely used as a turf grass (Bolaric et al., 2005a; 2005b; Wang et  
121 al., 2009), and for restoration and conservation seedings. Perennial ryegrass is naturally a  
122 diploid species ( $2n = 2x = 14$ ) (Bolaric et al., 2005b; Wang et al., 2009), but tetraploid ( $2n = 4x =$   
123 28) cultivars have also been developed (Nair, 2004). The grass has a two-locus self-

124 incompatibility system, which leads to an obligately outcrossing mating system (Cornish et al.  
125 1979). This mating system ensures outbreeding among individuals and high amounts of genetic  
126 diversity within naturally occurring populations (Bolaric et al., 2005a; 2005b; Wang et al., 2009).  
127 Many cultivars of perennial ryegrass however are derived from a limited pool of foundational  
128 clones; and such cultivars typically exhibit a limited amount of genetic variation (Guthridge et al.,  
129 2001).

130 Because perennial ryegrass is a cool season grass of temperate regions, it is generally  
131 considered to be sensitive to heat stress (Li et al., 2020); although heat tolerant cultivars have  
132 been developed (Wilkins, 1991). In addition, because the grass is widely cultivated and has high  
133 economic value, experiments assessing heat stress in perennial ryegrass, especially comparisons  
134 of heat tolerant and heat sensitive cultivars, have been conducted (e.g., Wehner and Watschke,  
135 1981; Jiang and Huang, 2001; Zhou and Abaraha, 2007; Wang et al., 2017; Sun et al., 2020; Li et  
136 al., 2020). For example, in a previous study we reported that a heat tolerant cultivar of perennial  
137 ryegrass (Yatugadake-24) exhibited significantly higher photosynthetic performance (i.e., they  
138 had higher Fv/Fm values) and lower leaf H<sub>2</sub>O<sub>2</sub> content, compared to a heat sensitive cultivar  
139 (Norlea) (Soliman et al., 2011). In another study, Soliman et al. (2012) exposed 25 diploid and  
140 tetraploid cultivars of perennial ryegrass to prolonged heat stress and found that tetraploid  
141 cultivars had lower H<sub>2</sub>O<sub>2</sub> content and experienced less oxidative stress than diploid cultivars.  
142 Taken together, these studies indicate considerable genetic variation in heat tolerance among  
143 perennial ryegrass cultivars and cytotypes. Yet, to the best of our knowledge, we are not aware  
144 of any assessment of the genetic basis of heat tolerance in perennial ryegrass.

145 In this study, we assessed the inheritance of heat tolerance in perennial ryegrass through a  
146 direct comparison of parental cultivars with their offspring, through progeny array analysis. This

147 was accomplished by crossing a heat tolerant cultivar of perennial ryegrass with a heat sensitive  
148 cultivar, to generate F1 progeny. Both parents and their progeny were then exposed to long-term  
149 heat stress, and their photosynthetic performance and leaf H<sub>2</sub>O<sub>2</sub> concentrations were measured.  
150 In addition, several leaf growth parameters were measured before the imposition of heat stress.  
151 The specific goals of this research were to, 1) quantify the level of heat tolerance in the two  
152 parental cultivars, 2) determine variation in heat tolerance among the F1 progeny, and 3)  
153 compare the level of heat tolerance of the parents with their F1 progeny to assess the inheritance  
154 of this complex and important phenotypic trait. Results of this study will improve our  
155 understanding of the genetic basis of heat tolerance in perennial ryegrass, assist in estimating the  
156 heritability of this trait, and aid in the identification and selection of plants with even higher  
157 levels of heat tolerance for use in plant breeding programs.

158

## 159 **MATERIALS AN METHODS**

### 160 **Plant material**

161 Two diploid perennial ryegrass cultivars were used as the parents in this study. Kangaroo Valley  
162 (strain K7) is a heat tolerant cultivar developed in New South Wales, Australia, that is well-  
163 suited to dry, hot regions ([Wilkens, 1991](#); [Blumenthal et al., 1996](#)) and Norlea (strain N4) is a  
164 heat sensitive cultivar developed in Canada ([Soliman et al., 2011](#); [2012](#)). Based on the breeding  
165 programs that developed them, both strains of the two cultivars exhibit limited genetic diversity  
166 ([Blumenthal et al., 1996](#); [Soliman et al., 2011](#)). Flowers of each of these two cultivars were  
167 crossed through hand-pollination, after they were emasculated. The Kangaroo Valley cultivar  
168 was always used as the pollen donor, and the Norlea cultivar always served as the maternal

169 parent. These crosses were conducted at the Yamanashi Dairy Experimental Station, Yamanashi,  
170 Japan. Seventy-two full-sib, F1 progeny derived from this cross were used in this experiment.

### 171 **Heat stress treatment**

172 Our heat stress experiment was conducted using the procedures described by [Soliman et al.](#)  
173 [\(2012\)](#). The seeds/seedlings of the two perennial ryegrass strains (K7 and N4) used in the heat  
174 stress experiments were not the same individuals used to generate the progeny array; but because  
175 these two strains have limited genetic diversity, seeds of these two cultivars are genetically  
176 uniform. Seeds of the two parental cultivars and the progeny were germinated on wet filter paper  
177 in petri dishes. The grass does not require any other treatments to achieve high rates of  
178 germination. Seedlings were transplanted into pots (two seeds per pot), 7.5 cm in diameter and 8  
179 cm deep, with a sandy loam potting soil containing 0.35 g of N, P<sub>2</sub>O<sub>5</sub>, and K<sub>2</sub>O for every  
180 kilogram of soil. The seedlings were grown in a controlled growth chamber with day/night  
181 temperatures of 23/16 °C, a 16h/8h day/night photoperiod (from 4:00-20:00 h), with photon flux  
182 density of 250  $\mu\text{mol m}^{-2} \text{s}^{-1}$ , and a constant relative humidity of 70%. Forty days after  
183 transplanting, all the plants were exposed to 30 °C for 3 days for acclimation to a higher  
184 temperature, after which the plants were exposed to heat treatments (36/30 °C, day/night  
185 temperatures) for 40 day. The heat stress experiment was set up in a randomized complete block  
186 design.

### 187 **Leaf growth traits**

188 Leaf growth traits were measured prior to the imposition of heat stress. For each seedling, the  
189 second fully matured leaf was selected for measurements. Specific leaf area (SLA) and its  
190 components were determined according to the method of [Witkowski and Lamont \(1991\)](#).  
191 Specific leaf area is calculated as the ratio of leaf area (LA) to leaf dry mass (LDM). Other leaf

192 measurements included leaf water content (LWC), leaf thickness (LT), and leaf density (LD).  
193 Before the start of the heat stress treatment, one mature leaf from each plant was harvested, its  
194 fresh weight was recorded, and it as immediately soaked in a 50 ml flask filled with water to  
195 perform measurements with leaves at full turgor. An image of each leaf was then digitally  
196 recorded using an optical scanner (D660U, Canon, Tokyo, Japan). The leaf area was calculated  
197 using Image J software (version 1.6, National Institutes of Health). The leaves were then oven  
198 dried at 80 °C for two days, and their dry weights were recorded. Leaf thickness was determined  
199 using microscopic observation of leaf transverse sections using MICROM (HM400R, Walldorf,  
200 Germany) as previously described (Soliman et al., 2012). Leaf density ( $\text{mg}/\text{cm}^3$ ), or dry matter  
201 concentration, was calculate by dividing leaf dry mass by leaf volume. Leaf volume was  
202 determined as the product of leaf area and mean leaf thickness.

### 203 **Photosynthetic performance**

204 Chlorophyll fluorescence ( $F_v/F_m$ ) values were measured before the initiation of heat acclimation  
205 and at 10-day intervals thereafter. Individual seedlings were maintained in the dark for 20 min  
206 for dark adaptation and then the minimum ( $F_0$ ) and maximal ( $F_m$ ) levels of fluorescence were  
207 measured three times for each individual using a portable photosynthesis measuring system (LI-  
208 6400, Li-Cor, Lincoln, Nebraska, USA).  $F_v/F_m$  provides an estimate of the maximum quantum  
209 yield of PSII (Butler, 1978; Zhou et al., 2015); where heat tolerant plants typically exhibit higher  
210  $F_v/F_m$  values (i.e., they have higher photosynthetic performance) than heat sensitive plants.

### 211 **Oxidative stress**

212 Heat sensitive plants experience greater oxidative stress than heat tolerant plants because plants  
213 that are heat sensitive (i.e., experiencing heat stress) produce higher concentrations of  $\text{H}_2\text{O}_2$  than  
214 heat tolerant plants. Hydrogen peroxide ( $\text{H}_2\text{O}_2$ ) concentration values were determined according

215 to the methods described by [Soliman et al. \(2011\)](#). There were two H<sub>2</sub>O<sub>2</sub> measurement periods;  
216 before the imposition of heat stress and at 40 days of exposure to heat stress. Hydrogen peroxide  
217 (H<sub>2</sub>O<sub>2</sub>) content of leaves was measured using a modified version of the ferrous ammonium  
218 sulphate/xylenol orange (eFOX) method described by [Cheeseman \(2006\)](#) and [Queval et al.](#)  
219 [\(2008\)](#). Leaf extracts were obtained by grinding 50 mg of leaf tissue, first in liquid nitrogen and  
220 then in 500 µL of 0.1 M potassium phosphate buffer (pH 6.5) containing 5 mM NaN<sub>3</sub>. Extracts  
221 were centrifuged at 10,000 rpm (8385 g) at 5°C for 5 min. For every 200 µL of the extract, 5 mL  
222 of the solution containing 250 µM ferrous ammonium sulphate, 100 µM sorbitol, 100 µM  
223 xylenol orange, 1% ethanol, and 25 mM H<sub>2</sub>SO<sub>4</sub> were added. The assay consisted of measuring  
224 the difference in absorbance between 550 nm and 800 nm, after 15 min, with a  
225 spectrophotometer.

## 226 **Statistical analyses**

227 Analysis of variance (ANOVA) was used to test for significance differences between the two  
228 parental cultivars and among the progeny for leaf growth traits; and among the progeny for  
229 Fv/Fm values and H<sub>2</sub>O<sub>2</sub> content, at different days of exposure to heat stress. Because the same  
230 plants were used to measure Fv/Fm values over time, and these data were not independent of  
231 each other, we used one-way repeated measures multivariate analysis of variance (MANOVA) to  
232 test whether the two cultivars were significantly different. We used a t-test to test for significant  
233 differences in the H<sub>2</sub>O<sub>2</sub> content between two cultivars before and at 40 days of heat stress. A  
234 random-effects regression model was used to assess the relationship between Fv/Fm values and  
235 H<sub>2</sub>O<sub>2</sub> content at 40 days after the imposition of heat stress, with Fv/Fm values and H<sub>2</sub>O<sub>2</sub> content  
236 as random variables. All statistical analysis were carried out using JMP (ver 4. SAS Institute,  
237 Cary, NC, USA).

238

## 239 **Results**

### 240 **Prior to heat stress treatment**

241 Before the imposition of heat stress, there were no statistically significant differences in  
242 chlorophyll fluorescence (Fv/Fm) values and hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) concentration values for  
243 the two parental cultivars (Figs. 1 and 2).

244 Prior to experiencing heat stress, the two parental cultivars did not exhibit significant  
245 differences for three of five leaf growth traits (Fig. 3). The two exceptions to this patter were leaf  
246 water content and leaf thickness. Conversely, significant variation was observed in all leaf traits  
247 among the progeny (Table 1), and they exhibited a normal distribution for all five leaf growth  
248 traits (Fig. 3). The majority of the 72 progeny had values of leaf growth traits beyond the values  
249 of the two parental cultivars. However, for leaf water content, 65% of the progeny had values  
250 that were intermediate to the values of the two parental cultivars (Fig. 3c). Seventy of 72 progeny  
251 had larger leaf area values compared to the two parents (Figure 3a). Most of the progeny  
252 generally had higher specific leaf area, thinner leaves, and higher leaf density compared to the  
253 two parents (Figures 3b, 3d and 3e, respectively). There was no relationship (no significant  
254 correlations) between leaf growth traits and Fv/Fm values and H<sub>2</sub>O<sub>2</sub> concentrations at 40 days  
255 after the imposition of heat stress (data not shown).

### 256 **Response to heat stress: parental cultivars**

257 The two parental cultivars did not exhibit statistically significant differences in Fv/Fm values for  
258 the first three measurement periods (0 d) and after the imposition of heat stress (10 d and 20 d).  
259 However, the two parents did show significant differences in their Fv/Fm values at 30 d and 40 d  
260 of heat stress, with the Kangaroo Valley cultivar having higher values (i.e., higher photosynthetic

261 performance) (Figs. 1 and 4a). After 40 d of to heat stress, both parental cultivars had higher  
262 H<sub>2</sub>O<sub>2</sub> values compared to before the imposition of stress. At 40 d of exposure to heat stress,  
263 Norlea, the heat-sensitive cultivar, had significantly higher H<sub>2</sub>O<sub>2</sub> content than the Kangaroo  
264 Valley cultivar (Figs. 2 and 4b). This result indicates that the Norlea cultivar experienced more  
265 oxidative stress under these conditions.

### 266 **Response to heat stress: progeny array**

267 Fv/Fm values for the 72 progeny were significantly different for all time periods measured (0 d  
268 to 40 d) (Table 2). Fv/Fm values of the progeny decreased gradually with increased duration of  
269 heat stress (Table 2), and a broad distribution of Fv/Fm values was observed at 40 d (Fig. 4a).  
270 Approximately 90% of the progeny had Fv/Fm values that were intermediate to the values of the  
271 two parental cultivars.

272 The progeny had statistically significant variation in H<sub>2</sub>O<sub>2</sub> values both without heat stress (at  
273 0 d) and at 40 days after the imposition of heat stress (at 40 d). Additionally, the progeny  
274 undergoing heat stress experienced an increase in their H<sub>2</sub>O<sub>2</sub> content (Table 2, Fig. 4b). At 40  
275 days after the imposition of heat stress, approximately 67% of the progeny had H<sub>2</sub>O<sub>2</sub> values that  
276 were equal to, or intermediate to, the two parental cultivars.

277 At 40 days of exposure to heat stress, the two cultivars and their progeny exhibited a  
278 significant inverse relationship ( $r = -0.54$ ) between Fv/Fm values and H<sub>2</sub>O<sub>2</sub> values (Fig. 5). Most  
279 of the data points for the progeny in Figure 5 cluster near the Kangaroo Valley cultivar, which  
280 served as the maternal parent in the cross that produced the progeny.

281

### 282 **Discussion**

283 Perennial ryegrass is one of the most common pasture and turf grasses in temperate climate  
284 regions around the globe (Bolaric et al., 2005a; 2005b; Wang et al., 2009). Because it is a cool-  
285 season grass, it is thought to be sensitive to heat stress (Li et al., 2020); however, plant breeders  
286 have also developed heat tolerant cultivars (Wilkins, 1991). In addition, because many strains of  
287 these cultivars are derived from a limited number of individuals, and possess limited genetic  
288 diversity (Guthridge et al., 2001), they can function similarly to inbred lines. These features of the two  
289 perennial ryegrass cultivars proved useful in designing our study. First, the Kangaroo Valley  
290 cultivar is heat tolerant and Norlea is heat sensitive; therefore, these two cultivars are genetically  
291 and phenotypically distinct. Second, different seeds of each of the two strains are genetically  
292 (and phenotypically) uniform, thus we could reliably use different seeds of each cultivar to  
293 generate the progeny and in the heat stress experiment.

294 The results of the current study are generally consistent with others that have assessed  
295 photosynthetic performance and oxidative stress with heat stress in heat tolerant and heat  
296 sensitive cultivars of perennial ryegrass (Wehner and Watschke, 1981; Jiang and Huang, 2001;  
297 Zhou and Abaraha, 2007; Soliman et al., 2011; 2012; Li et al., 2020). These results show that  
298 heat tolerant cultivars of perennial ryegrass had significantly higher photosynthetic performance  
299 (higher Fv/Fm values) and lower leaf H<sub>2</sub>O<sub>2</sub> content, compared to heat sensitive cultivars. At 40  
300 days of heat stress, approximately 90% of the progeny had Fv/Fm values that were intermediate  
301 to the values of the two parental cultivars and 67% of the progeny had H<sub>2</sub>O<sub>2</sub> concentrations  
302 intermediate to their two parents. Conversely, other members of this progeny array have  
303 phenotypic trait values beyond their two parental cultivars.

304 The phenotypic trait distribution for the five leaf growth traits, Fv/Fm values, and H<sub>2</sub>O<sub>2</sub>  
305 content for the 72 progeny is consistent with the distribution expected for traits that are

306 determined by multiple loci (i.e., they are quantitative genetic traits) (Falconer and Mackay,  
307 1996). The distribution for these phenotypic traits indicates considerable additive genetic  
308 variation within this progeny array, which resulted from crossing the Kangaroo Valley and  
309 Norlea cultivars. This variation was generated by genetic recombination during gamete  
310 formation by the parental plants. In addition, the clustering of many progeny data points near the  
311 Kangaroo Valley cultivar, which served as the paternal parent in the cross, may signal the role of  
312 dominance (the Kangaroo Valley cultivar possesses dominant alleles) or epistatic interactions in  
313 the expression of photosynthetic performance and leaf H<sub>2</sub>O<sub>2</sub> content in the progeny (Falconer  
314 and Mackay, 1996). Determining the relative contributions of additive genetic variation, and  
315 other genetic processes, in the expression of these two quantitative traits should be the focus of  
316 future research.

317 The photochemical efficiency of photosystem II (PSII), measured by chlorophyll  
318 fluorescence (Fv/Fm), is the most sensitive component associated with photosynthesis and it is  
319 used commonly to evaluate heat tolerance in plants (Maxwell and Johnson, 2000). Under  
320 elevated temperatures, ROS are produced through specific metabolic pathways such as  
321 photosynthesis and photorespiration (Queval et al., 2008). The generation of ROS results from  
322 disrupted balance between photochemical and biochemical reactions inhibiting the  
323 photosynthesis process (Wahid et al., 2007). Plants however have developed several mechanisms  
324 for tolerance to stress such as antioxidant enzymes and heat shock proteins.

325 The distribution of Fv/Fm and H<sub>2</sub>O<sub>2</sub> values among the progeny suggests the genetic variation  
326 for the genes responsible for heat tolerance. These genes control antioxidants activity and the  
327 formation of heat shock proteins, which in turn inhibit the formation of ROS and maintain  
328 membrane stability and thus increase photosynthetic efficiency, improve plant growth, and allow

329 plants to endure heat stress. These results suggest that the difference in heat tolerance shown by  
330 the 72 progeny analyzed in this study is closely associated with the ability to suppress oxidative  
331 stress. This is consistent with previous findings among cultivars of perennial ryegrass (Soliman  
332 et al., 2011, 2012).

333 Leaf growth traits also play important roles in plant acclimation to environmental stress  
334 (Terashima et al., 2018). We did not however detect a relationship between the leaf growth traits  
335 we measure prior to the imposition of heat stress and Fv/Fm values and H<sub>2</sub>O<sub>2</sub> content at 40 days  
336 after the imposition of heat stress. Results of the current study differ from those of our previous  
337 findings with other perennial ryegrass cultivars (Soliman et al., 2011), which showed significant  
338 relationships between leaf traits, especially leaf thickness, and ROS generation and heat  
339 tolerance. This discrepancy likely results from genetic difference of the parental cultivars used in  
340 the previous study. Clearly, heat tolerance is a complex phenotypic trait governed by many  
341 factors, not least of which is the genetic background of the plants (cultivars) being studied.

342

## 343 **Conclusions**

344 To the best of our knowledge, this study represents the first assessment of the genetic basis of  
345 heat tolerance in perennial ryegrass. This study combined physiological measurements (Fv/Fm  
346 and H<sub>2</sub>O<sub>2</sub> content) within a genetic framework (i.e., parent-offspring comparison) to assess the  
347 inheritance of heat tolerance in this grass. Based on the specific cross conducting in this study  
348 (the Kangaroo Valley and Norlea cultivars), our results indicate considerable additive genetic  
349 variation within this progeny array. This diversity could be used to improve heat tolerance in  
350 cultivars of perennial ryegrass using conventional plant breeding, and could also facilitate  
351 marker-assisted breeding and/or pave the way for characterizing the underlying genetic and

352 genomic factors which could be useful for developing plants with improved heat tolerance  
353 (Sreenivasulu et al., 2007; Barnabás et al. 2008; Tricker et al., 2018).

354

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358

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**Table 1** (on next page)

Table 1. Analysis of Variance for chlorophyll fluorescence (Fv/Fm) and hydrogen peroxide content ( $\text{H}_2\text{O}_2$ ,  $\mu\text{mol mgFW}^{-1}$ ) among 72 genotypes at different durations of continuous exposure to heat stress.

1

2

Table 1. Analysis of Variance (ANOVA) for leaf growth traits for the two parental cultivars and the 72 F1 progeny, prior to the imposition of heat stress.

Leaf growth traits	Parental cultivars			72 F1 progeny	
	Norlea	Kangaroo	F value	Range	F value
Leaf area (cm <sup>2</sup> )	1.72	1.81	0.07 <sup>ns</sup>	0.87 ~ 7.44	7.48 <sup>***</sup>
Specific leaf area (mm <sup>2</sup> mg <sup>-1</sup> )	26.96	24.43	1.01 <sup>ns</sup>	19.0 ~ 35.4	3.38 <sup>***</sup>
Leaf water content (%)	81.3	77.5	8.96 <sup>*</sup>	74.5 ~ 85.2	4.76 <sup>***</sup>
Leaf thickness (μm)	169	198	7.75 <sup>*</sup>	141 ~ 242	5.68 <sup>***</sup>
Leaf density (mg cm <sup>-3</sup> )	221	211	0.27 <sup>ns</sup>	153 ~ 328	7.93 <sup>***</sup>

\* and \*\*\* indicate the level of statistical significance at  $P < 0.05$  and  $P < 0.001$ , respectively.

<sup>ns</sup> indicates no statistical differences for the two parental cultivars for three leaf growth traits.

3

**Table 2** (on next page)

Table 2. Analysis of Variance for leaf traits between the two parents and among the 72 genotypes derived from them before exposure to heat stress.

1

Table 2. Analysis of Variance (ANOVA) for chlorophyll fluorescence (Fv/Fm) values and hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>, μmol mgFW<sup>-1</sup>) content among the 72 F1 progeny at different days of continuous exposure to heat stress.

Days of exposure	Range	F value
Chlorophyll fluorescence (Fv/Fm)		
0 day	0.762 ~ 0.807	2.71***
10 day	0.714 ~ 0.783	2.68***
20 day	0.603 ~ 0.778	3.01***
30 day	0.358 ~ 0.776	4.58***
40 day	0.483 ~ 0.767	25.43***
Hydrogen peroxide (H <sub>2</sub> O <sub>2</sub> )		
0 day	0.15 ~ 0.52	25.76***
40 day	0.32 ~ 1.74	27.95***

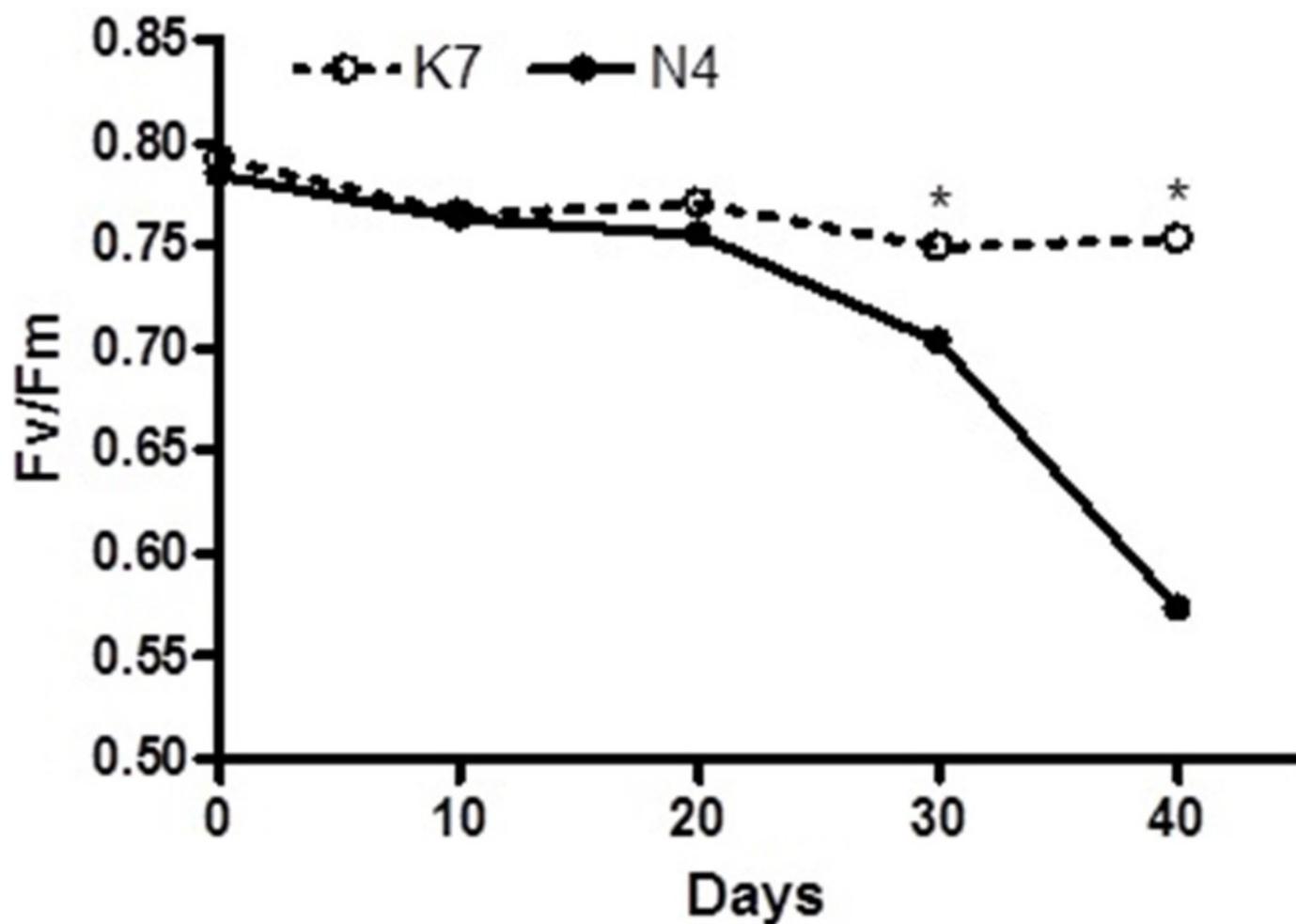
\*\*\* indicates the level of statistical significance at P < 0.001.

2

3

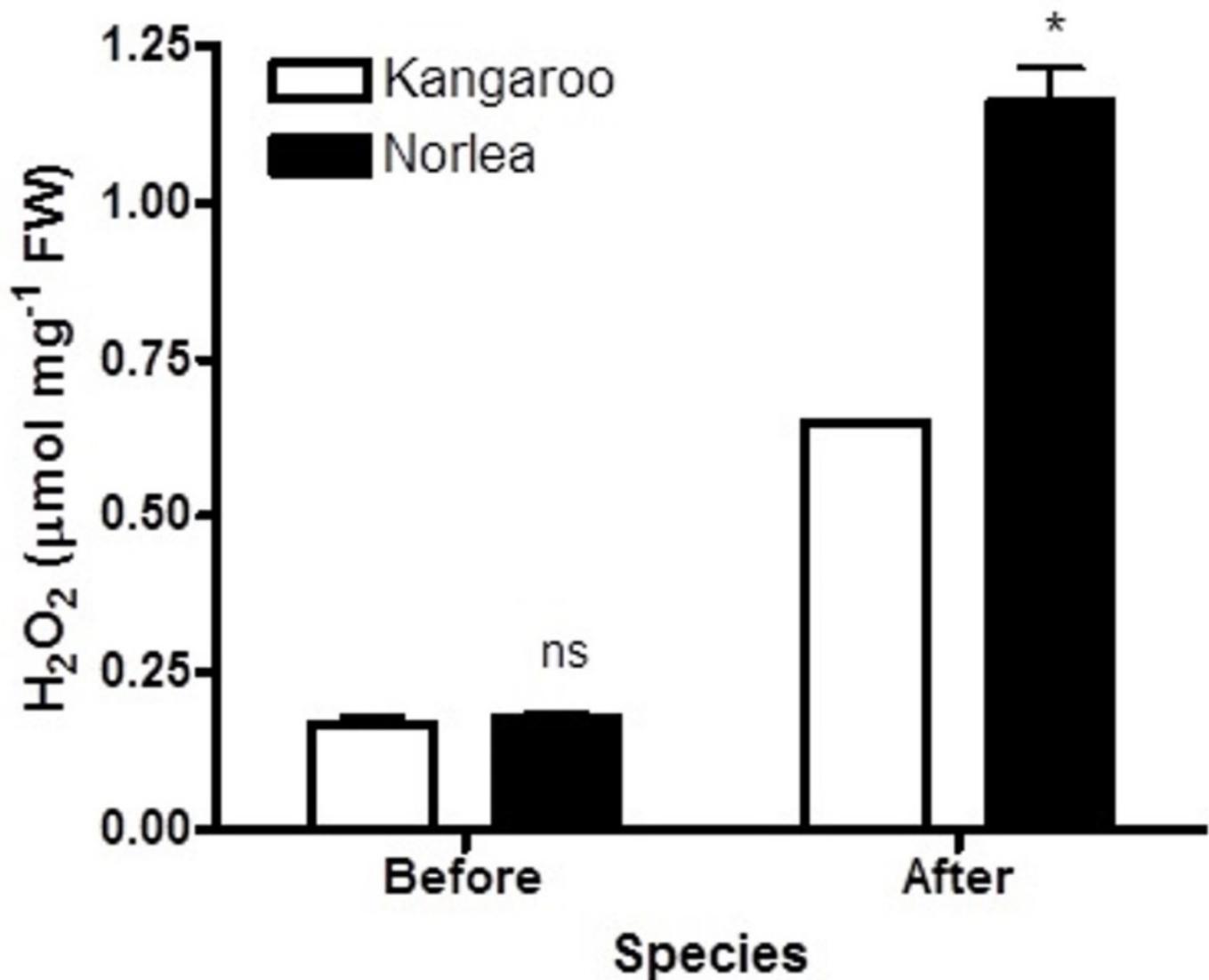
## Figure 1

Temporal changes in chlorophyll fluorescence ( $F_v/F_m$ ) values for the two parental cultivars, heat sensitive Norlea (N4) (●) and heat tolerant Kangaroo Valley (K7) (○), after imposition of heat stress. \* represents the level of statistical significance at P



## Figure 2

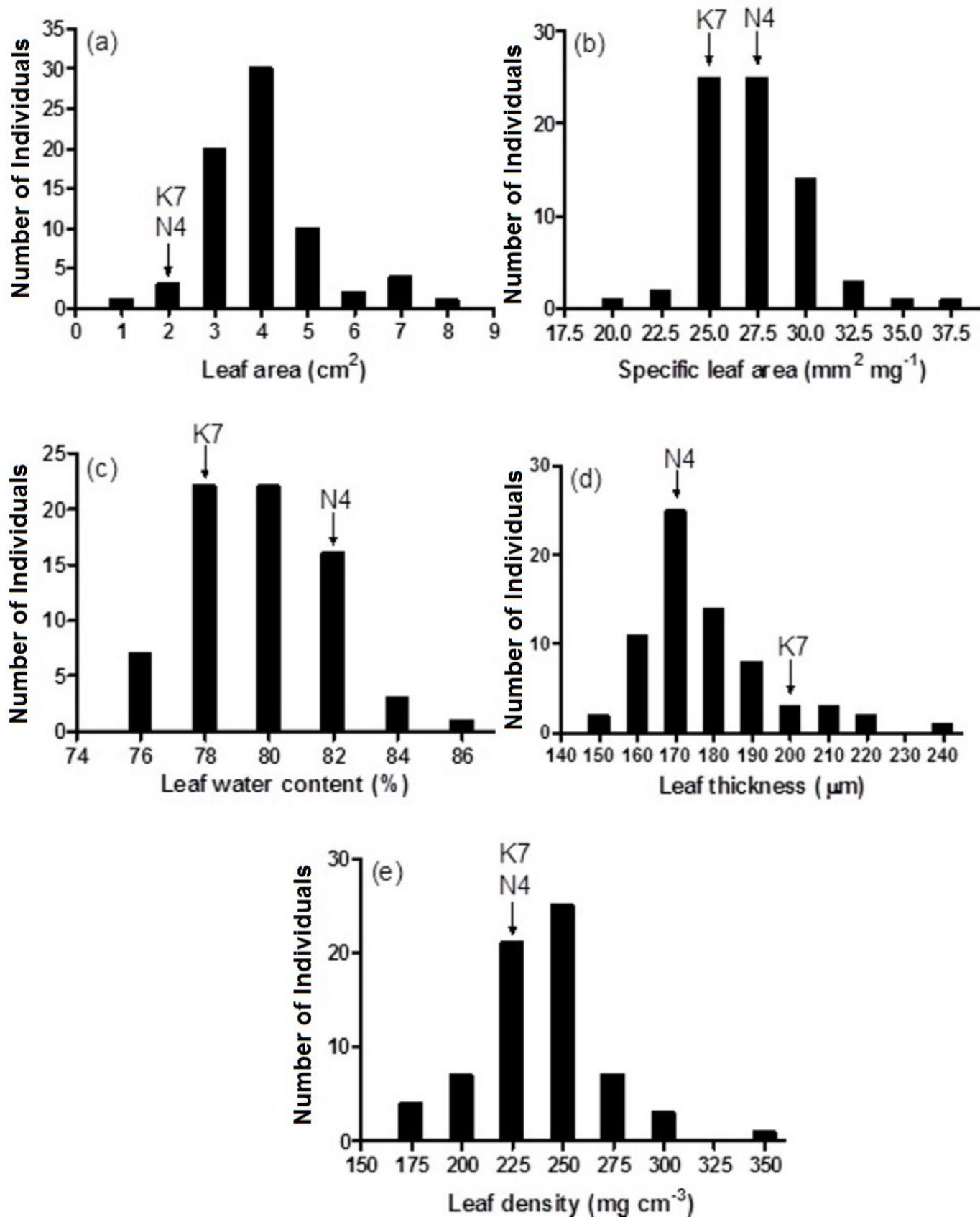
Hydrogen peroxide ( $H_2O_2$ ) content before and after the imposition of heat stress in the two parental cultivars; heat sensitive Norlea (N4) and heat tolerant Kangaroo Valley (K7). \* represents the level of statistical significance at  $P <$



\* represents the difference at 0.1%

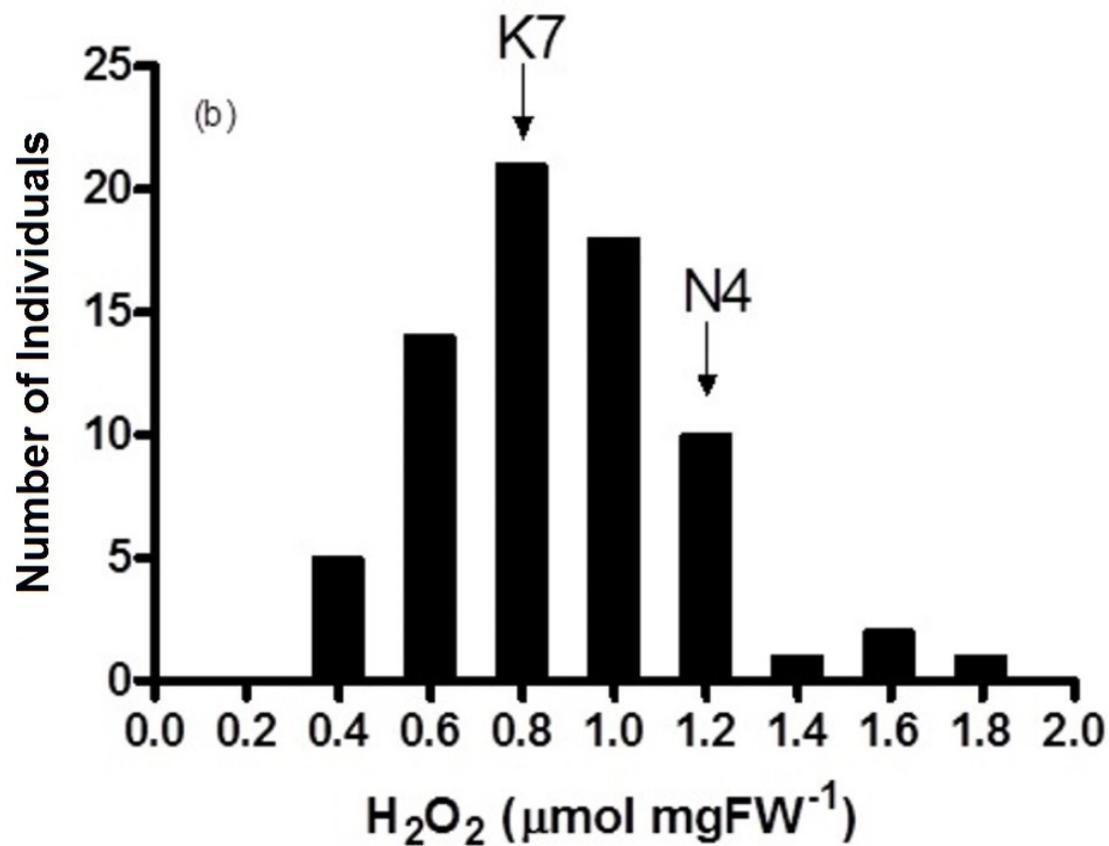
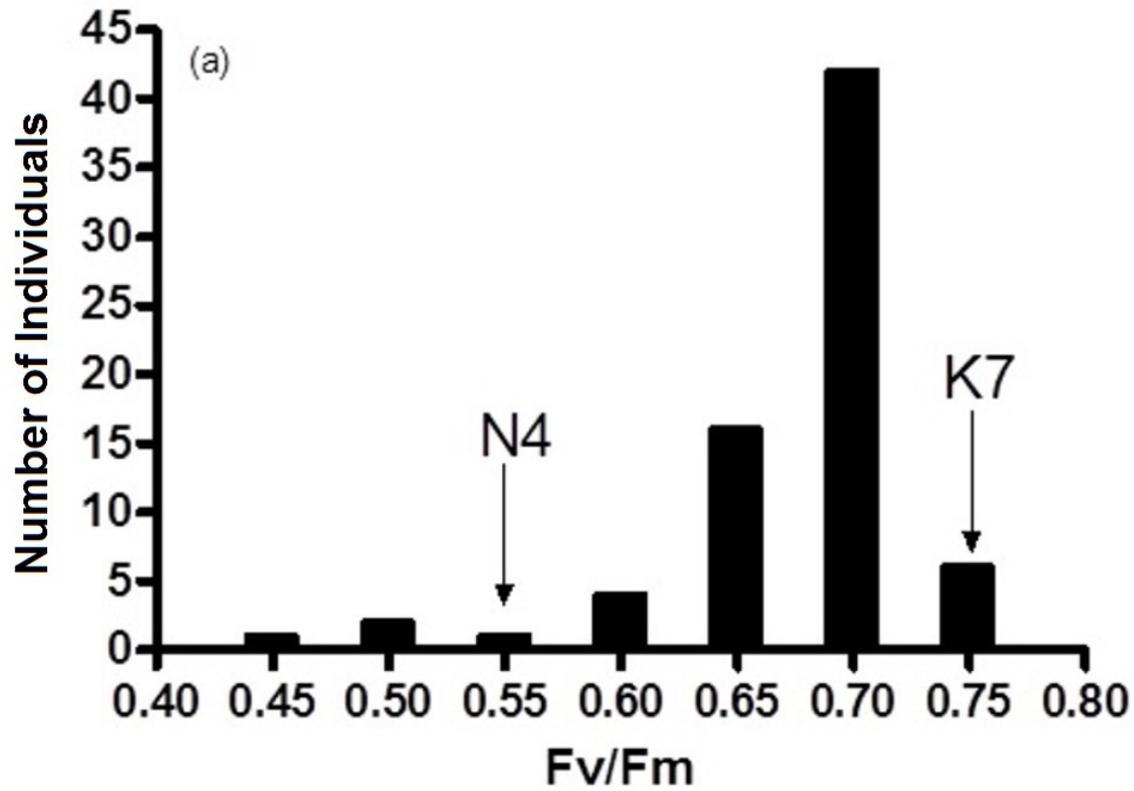
## Figure 3

Frequency distributions of leaf area (a), specific leaf area (b), leaf water content (c), leaf thickness (d) and leaf density (e) for the 72 F1 progeny and the two parental cultivars, heat tolerant Kangaroo Valley (K7) and heat sensitive Norlea (N4), prio



## Figure 4

Frequency distribution of chlorophyll fluorescence (Fv/Fm) values (a) and hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) content (b) for the 72 F1 progeny and the two parental cultivars, heat tolerant Kangaroo Valley (K7) and heat sensitive Norlea (N4), at



## Figure 5

Correlation between chlorophyll fluorescence ( $F_v/F_m$ ) values and hydrogen peroxide ( $H_2O_2$ ) content for the 72 F1 progeny and the two parental cultivars, heat tolerant Kangaroo Valley (K7) and heat sensitive Norlea (N4), at 40 days of exp

