



Examining Genetic Variation in Maize Inbreds and Mapping Oxidative Stress Response QTL in B73-Mo17 Nearly Isogenic Lines

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Specialty section:

This article was submitted to
Climate-Smart Food Systems,
a section of the journal
Frontiers in Sustainable Food Systems

Received: 10 April 2019

Accepted: 17 June 2019

Published: 04 July 2019

Citation:

Sorgini CA, Barrios-Perez I, Brown PJ
and Ainsworth EA (2019) Examining
Genetic Variation in Maize Inbreds and
Mapping Oxidative Stress Response
QTL in B73-Mo17 Nearly Isogenic
Lines. *Front. Sustain. Food Syst.* 3:51.
doi: 10.3389/fsufs.2019.00051

Screening crop plants under elevated ozone concentrations ([O₃]) is a pre-requisite for identification of tolerant lines, but few studies have mapped maize responses to elevated [O₃]. B73-Mo17 nearly isogenic lines (NILs) were screened in the field under ambient (~40 ppb) and elevated (~100 ppb) [O₃] at the Free Air gas Concentration Enrichment (FACE) research facility in Champaign, IL to identify maize leaf damage QTL associated with variation in O₃-induced oxidative stress response. In Mo17 NILs, a significant leaf damage QTL was identified at 161Mb on chromosome 2. To assess the feasibility of high-throughput phenotyping and fine mapping of early season O₃ leaf damage QTL, a subset of the nested association mapping (NAM) founder lines were screened in a growth chamber experiment under ambient and elevated [O₃]. Results showed that elevated [O₃] decreased the number of green leaves while increasing the number of lesioned and dead leaves. Most lines showed the same general response to elevated [O₃], but the degree of damage varied among lines. Next, tolerant and sensitive B73-Mo17 NILs identified from the FACE study, and hybrid crosses of the identified NILs with Mo17 (*n* = 20) were grown under elevated O₃ (~150 ppb) in growth chambers (*n* = 7). In the chambers, O₃-sensitive lines could be distinguished from tolerant lines based on leaf lesions, but there was not a continuous degree of damage like that seen in the field. This research identified a repeatable O₃-induced leaf damage QTL and developed populations and markers that can be used in future growth chamber fine mapping experiments. These results demonstrate the feasibility of high-throughput phenotyping and fine mapping of O₃ leaf damage QTL in a controlled environment.

Keywords: maize, tropospheric ozone (O₃), FACE technology, QTL mapping, global climate change

INTRODUCTION

Tropospheric O₃ is one of the most important environmental pollutants and is estimated to cost billions of dollars in global crop losses (Van Dingenen et al., 2009; Avnery et al., 2011; McGrath et al., 2015; Ainsworth, 2017; Mills et al., 2018). A majority of tropospheric O₃ comes from anthropogenic emissions. Tropospheric O₃ is a direct driver of global warming and has indirect negative effects

on plant production. O₃ has been shown to have a negative effect on yield and quality traits of crop plants such as soybean and rice (Betzberger et al., 2010; Frei, 2015). However, there have been fewer studies that have investigated the effects of elevated O₃ on C₄ plants (Leisner and Ainsworth, 2012). Maize is one of the world's primary agricultural commodities for food, fodder, and fuel (FAO, 2018). The global demand for maize crop production is increasing exponentially (Kay et al., 2013), and it is estimated that by 2050 agricultural commodities will need to sustain more than nine billion people (FAO, 2018). Concurrently, it is projected that by 2050 tropospheric O₃ concentrations will increase (Pachauri et al., 2014). Therefore, understanding how maize is affected by O₃-induced oxidative stress will contribute to improving current and future maize crop productivity.

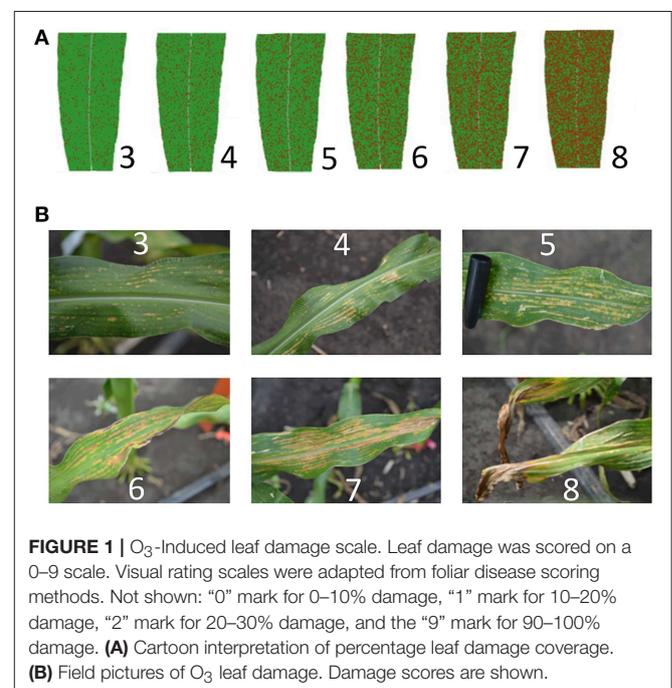
O₃ causes damage to plants when it enters leaves through the stomata and forms other reactive oxygen species (ROS; Black et al., 2000). Plant response to oxidative stress involves the creation of ROS stress and its interaction with phytohormones, Ca²⁺, and MAPK signal cascades. There appears to be significant overlap between O₃ response and pathogen response pathways in plants (Perez and Brown, 2014). High [O₃] mimics oxidative bursts generated by early signal pathways that regulate plant hypersensitive response (Rao and Davis, 1999; Rao et al., 2000). Secondary ROS bursts activate the expression of defense genes and the ethylene, salicylic acid, and jasmonic acid signal pathways (Black et al., 2000; Perez and Brown, 2014). The cost associated with detoxification of O₃ often increases respiration, which along with a reduction in photosynthesis at elevated [O₃], imparts a substantial C cost to plants (Ainsworth et al., 2012).

Amongst the range of effects caused by elevated [O₃], visible leaf damage symptoms are considered an important indicator of O₃ injury (Miller et al., 1989). O₃-induced leaf damage traits have been reported for the past 35 years from countries all around the world (Krupa et al., 2001). Both acute and chronic O₃ exposure induce oxidative stress due to the production of ROS in the apoplast (Frei, 2015). At high O₃ concentrations, this production of ROS can result in cell death and necrotic symptoms (Rao and Davis, 2001; Kangasjarvi et al., 2005). Chronic injury is normally induced by long-term, lower O₃ concentrations, and develops slowly over days to weeks. Chronic injury is characterized by chlorosis, stippling, necrosis, leaf edge yellowing, and premature senescence (Brace et al., 1999). A concern in diagnosing leaf damage is the ability to distinguish O₃ symptoms from a wide range of potential symptoms caused by other agents. Therefore, foliar symptoms are best identified through a systematic survey. Assessing O₃-induced leaf damage is important because it is often correlated with a decreased carbon fixation and water use efficiency (Rao and Davis, 2001; Kangasjarvi et al., 2005) and is much easier to score than physiological measurements that require specialized equipment.

Free Air gas Concentration Enrichment (FACE) facilities allow for the investigation into predicted climate change scenarios. The power of FACE facilities is in the ability to study how changes in atmospheric gasses alter plant growth in a field setting, with treatment application having only a minimal effect on other abiotic and biotic factors. FACE technologies have been adapted to enrich O₃ to study plant responses in field settings

(Morgan et al., 2004; Tang et al., 2011). A potential problem with FACE systems is their limited size, making it difficult to screen large populations needed for modern genetic analysis. Previous work has shown that the SoyFACE research facility at the University of Illinois Urbana-Champaign can be successfully used to screen crops for O₃ tolerance and sensitivity (Ainsworth et al., 2014; Yendrek C. R. et al., 2017). Using modern screening SoyFACE studies have shown genetic variation in numerous traits for numerous species. The most recent research in crop species response to elevated O₃ has focused on identifying physiological variation and/or yield traits in soybean (Betzberger et al., 2010), rice (Shi et al., 2009), and wheat (Zhu et al., 2011). Identifying intraspecific variation for oxidative stress tolerance is an important pre-breeding step (Ainsworth, 2017). However, the next step is to use genetically structured populations, such as near isogenic lines, in FACE experiments to perform genetic analysis of O₃ response. Genetic mapping studies have identified leaf damage QTL for Arabidopsis (Brosché et al., 2010), poplar (Street et al., 2011), rice (Kim et al., 2004; Frei et al., 2008; Frei, 2015; Tsukahara et al., 2013), and soybean (Burton et al., 2016), but not for maize.

Building on previous, unpublished results suggesting B73 and Mo17 differed in O₃ sensitivity, B73-Mo17 nearly isogenic lines (NILs) were screened in the field under ambient (~40 ppb) and elevated (~100 ppb) [O₃] at the FACE research facility in Champaign, IL to identify maize leaf damage QTL associated with variation in oxidative stress response. The aims of this study were to (i) assess variation in the sensitivity of maize to elevated [O₃] using nearly isogenic lines, (ii) identify O₃-induced leaf damage QTL(s) and, (iii) develop populations and marker tools for fine mapping to confer O₃ tolerance and/or verify sensitive lines. Additionally, the feasibility of high-throughput



phenotyping and fine mapping of early season O₃ damage QTL were tested with a controlled environment experiment. Two hypotheses were tested: (i) maize exposure to elevated [O₃] (~150 ppb) in a growth chamber will result in an abiotic stress response, which will accelerate senescence; and, (ii) O₃ response in the growth chamber will correlate to response measured in the field.

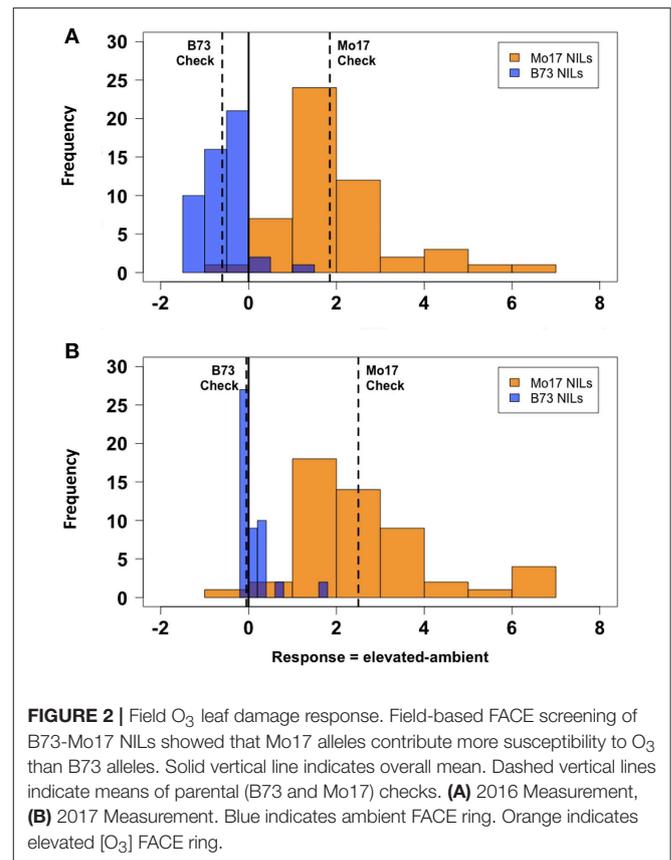
MATERIALS AND METHODS

Field NIL Experiment

One hundred B73-Mo17 NILs were screened in the field under ambient (~40 ppb) and elevated (~100 ppb) [O₃] at the FACE research facility in Champaign, IL. Fifty B73 NILs ($n = 4$) with Mo17 introgressions into a B73 background and 50 Mo17 NILs ($n = 4$) with B73 introgressions into a Mo17 background developed by Eichten et al. (2011) were planted in a single-row 1.65 m plot in four elevated [O₃] and four ambient rings in 2016 and 2017. B73 and Mo17 were grown as checks and replicated five times within each ring for a total of 40 plots across the experiment. Each NIL was present one time within each ring for a total of eight replications across the experiment. Air enriched with O₃ was delivered to the experimental rings with FACE technology, as described in Yendrek C. et al. (2017). Maize was exposed to elevated [O₃] (100 ppb) for 8 h each day from shortly after emergence until physiological maturity. When the fumigation system was operating, O₃ concentrations were within 20% of the 100 ppb target concentration for 81% of the time.

O₃-induced leaf damage was scored on a 0–9 scale in two successive years. A foliar disease scale was modified for leaf damage susceptibility in maize (Figure 1); with zero representing 0–10% of the leaf area having damage and nine representing 90–100% of the leaf area having damage. Leaf damage was scored on eight plants per plot and the average was used for analysis. Plot averages for each FACE ring were used in the analysis for each NIL genotype. In 2016 and 2017, 43 days after planting (DAP) measurements were taken on the 5th true leaf. Damage scores were collected independently by two scientists and compared for reliability.

QTL analysis was completed using stepwise regression with the *lm* function in R (R Core Team, 2015). Three models with different random effects were tested and best fit chosen by the Akaike information criterion (AIC). The final model included random effects for ring, ring set by ring interaction (ringset:ring), and genotype. “Ringset” is the cardinal direction location of the plot in each ring. This model was used to analyze the data in separate genotypes, in separate environments, and in both genotypes. First, the ozone response (Elevated leaf damage—Ambient leaf damage) was calculated. Mo17 NILs were analyzed separately from B73 NILs. Then, an additional QTL analysis was completed in separate environments (elevated or ambient). Finally, the analysis was run analyzing B73 and Mo17 NILs together with recurrent parent (“RP”) as a covariate in the model. Significance thresholds were determined by using 200 permutations and an alpha of 0.05.



Growth Chamber Experiments

A subset of nested association mapping (NAM) founder lines (CML322, CML333, Ki3, M37W, Mo18W, MS71, NC358, and P39) were grown under ambient and elevated [O₃] (~150 ppb) in growth chambers ($n = 4$) (Environmental Growth Chamber, GC Series, 1.4 m² growing area, Chagrin Falls, OH, USA). Ozone was produced by a variable output UV-C light bulb ballast (HVAC 560 ozone generator, Crystal Air, Langley, Canada), and controlled with a custom multiport sampling system. The subset of NAM founder lines was selected based on previous results (Yendrek C. R. et al., 2017) that showed in elevated [O₃] leaf senescence of inbred and hybrid maize was genotype-specific. The NAM founder lines were selected to maximize diversity from maize inbreds. The subset selected contained tropical lines (CML322, CML333, Ki3), a high-carotenoid line (M37W), temperate lines (Mo18W, MS71, NC358), and a popcorn (P39). Each chamber was setup in a 10 × 5 ($n = 50$ plants per chamber) layout, as an incomplete block design with five blocks and 10 plants per block. Each block had a B73 check such that B73 was replicated 10 times per chamber for a total of 80 pots across the experiment. The selected NAM founder lines were replicated five times per a chamber for a total of 40 pots across the experiment. Genotype location was randomized for ambient and elevated [O₃] chamber pair (Supplemental Figure 1). Growth conditions included SunGrow[®] professional LC1 growing mix soil, osymocote[®] all-purpose plant fertilizer, and daily hand

watering. Chambers were set to maintain constant light ($700 \mu\text{mol m}^{-2} \text{s}^{-1}$; 15 h day), temperature (25°C day, 21°C night), and relative humidity (60%). Ozone was produced with a variable output UV-C light bulb and ballast (HVAC 560 ozone generator, Crystal Air, Langley, Canada) for a setpoint of 150 ppb for 9 h per day. Ozone concentrations within each chamber were monitored with an O_3 analyzer (Thermo Electron 49i, Thermo Scientific,

Waltham, MA, USA) and controlled using a custom multiport sampling system.

Total leaf number, green leaf number, lesioned leaf number, and dead leaf number phenotypic measurements were completed at 21 and 32 DAP. Additional phenotypic measurements were completed 32 DAP and included height of the main stalk to the whorl (“height”), tiller number, and the sum of all tiller

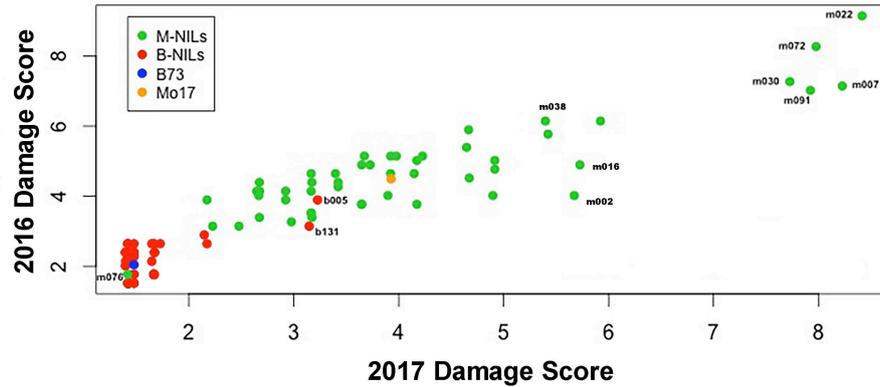


FIGURE 3 | Field O_3 leaf damage correlation. Leaf damage scores from the field in 2016 and 2017 had a strongly significant correlation ($r = 0.93$). Five sensitive Mo17 NILs (m007, m022, m030, m072, and m091), one tolerant Mo17 NIL (m076), and two sensitive B73 NILs (b005 and b131) were identified and consistent across years.

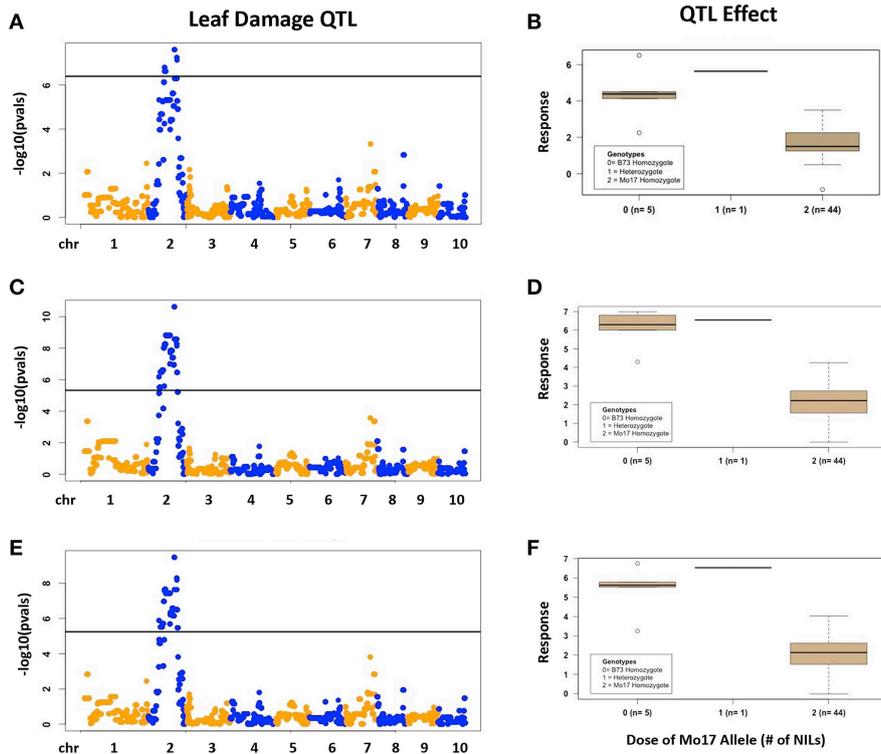


FIGURE 4 | Field O_3 Leaf Damage QTL in Mo17 NILs. Results are shown for Mo17 NILs; (A,B) 2016, (C,D) 2017, (E,F) 2016 and 2017 combined data results. (Left) QTL mapping results. (Right) Boxplots showing phenotypic distributions for B73-allele homozygotes (0), heterozygotes (1), and Mo17-allele homozygotes (2) at the most significant marker, which is the same in (A–C).

lengths and main stalk (“total height”). Green leaf number and lesioned leaf number were scored to capture different potential response mechanisms. In the field, some genotypes put leaves on more rapidly in elevated ozone, and so we counted green leaf number to determine if this trait was consistent. Lesioned leaf number informs us how many leaves are showing ozone sensitivity. Statistical analysis was performed on mean trait values for each genotype within a chamber. Independent *T*-tests were performed on genotype by trait:treatment. Significance was adjusted using the Bonferroni correction for multiple testing. Linear mixed effect modeling was completed using the Lme4 R package (Bates et al., 2015), using a model with: treatment as a fixed effect, chamber within treatment as a random effect, block within chamber within treatment as random effect, genotype as a fixed effect, and genotype*treatment as a fixed effect. The mixed model *p*-values were calculated using normal distribution approximation which assumes infinite degrees of freedom. The significance threshold was set at *p* < 0.05 for all analyses.

Selected tolerant (m076) and sensitive (m002, m072, m007, m030, m091, m016, m038, b131, b005) NILs plus their hybrid crosses with Mo17 were grown under elevated O₃ (~150 ppb) in growth chambers (*n* = 7), with an incomplete block design. Each chamber was setup in a 4 × 8 (32 plants per chamber) layout to allow plants to grow without overcrowding through the 6th leaf stage. Each chamber contained four blocks with eight plants per block. Each block had B73, Mo17, and B73 × Mo17 checks, so each check was replicated four times within a chamber. Selected NIL lines and hybrids were screened in each chamber for a total of seven plants in the experiment. Within a chamber plant location was randomized (Supplemental Figure 2). Growth conditions included SunGrow[®] professional LC1 growing mix

soil, osymocote[®] all-purpose plant fertilizer, and daily hand watering. Chambers were set to maintain constant light (700 μmol m⁻² s⁻¹; 15 h day), temperature (25°C day, 21°C night), relative humidity (60%), and elevated [O₃] (150 ppb) for 9 h per day.

O₃-induced leaf damage was scored on a 0–9 scale as described above, on the 5th and 6th true leaf at 32 DAP. Chamber average damage scores were recorded for each NIL, hybrid, and check. Leaf damage scores were collected independently by two scientists and compared for reliability. Additionally, each leaf of each plant was photographed. Growth chamber values for leaf damage were compared to field values scored in the elevated [O₃] plots using a 2 parameter logarithmic function ($y = y_0 + \alpha \ln x$). The data were assessed with the Shapiro Wilk normality test and QQ plot analysis using the 5th leaf damage scores from the chamber, the 6th leaf damage scores from the chamber, the combined 5th and 6th leaf (“leaf_variable”) damage scores, and the field leaf damage scored on 5th leaf in elevated [O₃]. To determine best fits for the data distributions, three models were tested and the best was chosen using AIC. The final models included random effects for leaf_variable, chamber, block:chamber, and genotype. All calculations and analysis were completed in R (R Core Team, 2015).

RESULTS

Field NIL Experiment

There was considerable genetic variation within the NILs in the response to elevated [O₃] (Figure 2). Mo17 was more sensitive than B73, and some Mo17 NILs were much more sensitive than Mo17 (Figure 2). Leaf damage scores from the field in 2016 and 2017 had a strong, significant correlation (*r* = 0.93, Figure 3). Five sensitive Mo17 NILs (m007, m022, m030, m072, and m091), two sensitive B73 NILs (b005, b131), and one tolerant Mo17 NIL (m076) were consistent across years (Figure 3). In Mo17 NILs, a repeatable significant QTL was identified on chromosome 2 at 161Mb (Figures 4A,C,E and Table 1). Interestingly, B73 introgressions into Mo17 in this region made NILs more sensitive to elevated [O₃] (Figures 4B,D,F). Sensitive Mo17 NILs had on average five total introgression regions. All five sensitive NILs shared a common introgression on chromosome 2 at 161Mb, and the left-hand boundaries of the LOD drop off support intervals (Supplemental Table 1) for this QTL cross the centromere. Sensitive B73 NILs (b005 and b131) had two shared introgressions: one on chromosome 5 that is 1.86Mb and another on chromosome 6 that is 7.95 Mb. Resistant NIL m076 has six small homozygous introgressions on four chromosomes.

NAM Founder Lines Subset Growth Chamber Experiment

Maize plant exposure to elevated [O₃] in a growth chamber setting resulted in an abiotic stress response that was distinguishable by leaf trait phenotyping. In general, independent *T*-tests by trait for each genotype show that elevated [O₃] in the growth chambers decreased the number of green leaves while increasing the number of lesioned and dead leaves

TABLE 1 | Field stepwise regression QTL linkage mapping results.

| Data values | Geno effect | Collection year | Step1 marker |
|-----------------|-------------|-----------------|--------------|
| Mo17 NIL | Response | 2016 | 1782*** |
| Mo17 NIL | Response | 2017 | 1782*** |
| Mo17 NIL | Response | 2016 + 2017 | 1782*** |
| Mo17 NIL | ele | 2016 | 1782*** |
| Mo17 NIL | ele | 2017 | 1782*** |
| Mo17 NIL | ele | 2016 + 2017 | 1782*** |
| Mo17 & B73 NILs | Response | 2016 | 1782*** |
| Mo17 & B73 NILs | Response | 2017 | 1782*** |
| Mo17 & B73 NILs | Response | 2016 + 2017 | 1782*** |
| Mo17 & B73 NILs | ele | 2016 | 1782*** |
| Mo17 & B73 NILs | ele | 2017 | 1782*** |
| Mo17 & B73 NILs | ele | 2016 + 2017 | 1782*** |

The analysis, which used the best fit linear effects model determined by AIC, identified the same QTL for separate genotypes, separate environments, and in both genotypes. First, Mo17 NILs were analyzed separate from B73 NILs. Then, B73 and Mo17 NILs analyzed together with recurrent parent (RP) as a covariate. Threshold significance was determined by using 200 permutations, alpha of 0.05. Geno effect column indicates if the model was run as a response QTL or separately in the elevated (“ele”) environment. ****p* < 0.001, no significant QTL were identified in the ambient environment alone. Marker # 1782 is located on chromosome 2, 160,938,561.9bp, 135.22cM (AGPv2).

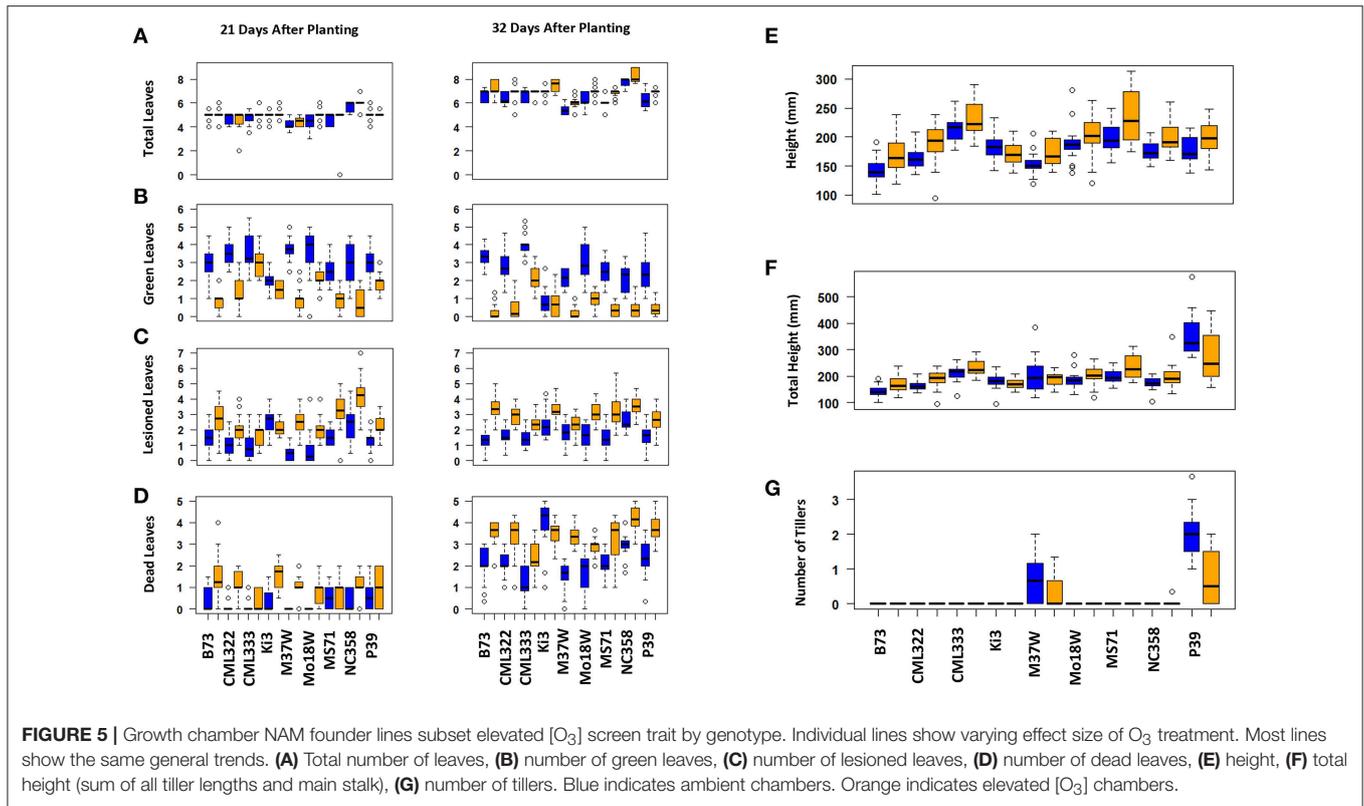


TABLE 2 | Growth chamber NAM founder lines subset elevated $[O_3]$ screen independent *T*-tests trait by genotype.

| | Total leaves | | Green leaves | | Lesioned leaves | | Dead leaves | | Height | Tillers | Total height |
|--------|--------------|----------|--------------|----------|-----------------|----------|-------------|----------|----------|----------|--------------|
| | 21 DAP | 32 DAP | 21 DAP | 32 DAP | 21 DAP | 32 DAP | 21 DAP | 32 DAP | 32 DAP | 32 DAP | 32 DAP |
| B73 | 4.41E-02 | 6.12E-06 | 1.98E-15 | 1.76E-15 | 9.34E-09 | 1.98E-15 | 8.83E-10 | 1.76E-15 | 1.27E-04 | NA | 1.27E-04 |
| CML322 | ns | ns | 4.62E-10 | 2.50E-10 | 6.11E-04 | 1.51E-07 | 2.48E-06 | 7.39E-06 | 3.27E-02 | NA | 3.27E-02 |
| CML333 | ns | ns | ns | 1.15E-10 | 7.55E-02 | 1.10E-05 | ns | 4.20E-05 | ns | NA | ns |
| Ki3 | ns | 3.77E-04 | 1.68E-02 | ns | ns | 7.75E-04 | 2.50E-07 | ns | ns | NA | ns |
| M37W | ns | 1.52E-02 | 1.98E-15 | 3.37E-14 | 2.63E-10 | 5.26E-02 | 1.68E-06 | 1.58E-10 | ns | 1.70E-02 | ns |
| Mo18W | 5.97E-03 | 6.10E-04 | 1.16E-03 | 5.77E-07 | 1.14E-03 | 2.74E-06 | 1.93E-05 | 2.24E-06 | ns | NA | ns |
| MS71 | ns | 7.42E-08 | 6.99E-08 | 4.50E-12 | 2.48E-06 | 6.52E-06 | ns | 4.00E-03 | 2.82E-02 | NA | 2.82E-02 |
| NC358 | 2.42E-02 | 1.80E-02 | 1.81E-06 | 2.40E-08 | 1.07E-04 | 1.08E-03 | 3.22E-03 | 1.99E-06 | 2.89E-02 | NA | 2.89E-02 |
| P39 | ns | 1.18E-02 | 4.92E-06 | 3.05E-07 | 5.42E-04 | 4.94E-04 | ns | 1.37E-07 | ns | 3.34E-07 | 2.40E-02 |

Maize plant exposure to elevated O_3 in a growth chamber setting resulted in an abiotic stress response that was distinguishable by leaf trait phenotyping. Direction of effect is indicated by color coding. Blue indicates significant increase of trait values in ambient conditions. Orange indicates significant increase of trait values in elevated conditions. Total height is the sum of all tiller lengths and main stalk. Dark shading ($p < 0.01$) and light shading ($p < 0.05$). Significance adjusted for multiple testing using Bonferroni. ns, not significant, NA, no tillers available to measure.

(Figure 5 and Table 2). Most NAM founder lines showed the same general trends for leaf traits, however individual lines showed varying responses to elevated O_3 treatment (Figure 5 and Table 2). Height response was variable; lines B73, CML322, MS71, and NC358 were significantly ($p < 0.05$) affected by O_3 treatment, but lines CML333, Ki3, M37W, and Mo18W were not (Table 2). Line P39 tiller production was significantly ($p < 0.05$) reduced in the elevated O_3 environment (Table 2). The growth chamber experiment showed that B73 grown

under ~ 150 ppb O_3 for 3–4 weeks can clearly be differentiated from ambient-grown B73 based on leaf lesion types (Table 2). Linear mixed-effects modeling shows that NAM founder lines B73 and MS71 were significantly affected ($p < 0.05$) by O_3 treatment in all three traits analyzed (Table 3). NAM founder lines CML322, M37W, Mo18W, Ms71, and NC358 were significantly affected ($p < 0.01$) by O_3 treatment for traits green leaf number and lesioned leaf number but not height (Table 3).

TABLE 3 | Growth chamber NAM founder lines subset *P*-values from Lme4 modeling.

| | Green 21 DAP | Lesion 21 DAP | Height |
|--------|-----------------|------------------|----------|
| B73 | 0.00E+00 | 2.52E-05 | 4.68E-02 |
| CML322 | 2.22E-16 | 1.48E-03 | ns |
| CML333 | 1.96E-02 | 3.41E-02 | ns |
| Ki3 | 5.18E-02 | ns | ns |
| M37W | 0.00E+00 | 8.69E-10 | ns |
| Mo18W | 7.22E-08 | 1.10E-04 | ns |
| MS71 | 4.83E-10 | 5.86E-07 | 2.09E-02 |
| NC358 | 2.22E-16 | 2.64E-08 | ns |
| P39 | 1.88E-06 | 5.10E-03 | ns |

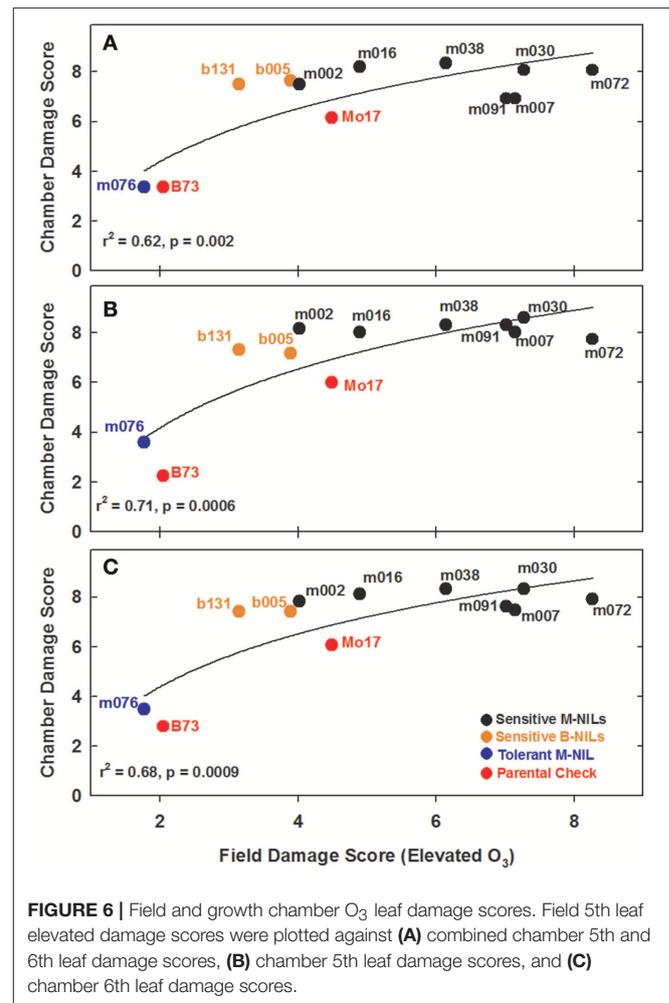
Lme4 estimated *p*-values show that NAM founder lines B73 and MS71 were significantly affected by O₃ treatment in all three traits analyzed ($p < 0.05$). NAM founder lines CML322, M37W, Mo18W, NC358, and P39 were significantly affected ($p < 0.01$) by O₃ treatment for traits green leaf number and lesioned leaf number but not height. Direction of effect indicated by color coding. Orange indicates significant increase of trait values in elevated conditions. Blue indicates significant increase of trait values in ambient conditions. Dark shading ($p < 0.01$), light shading ($p < 0.05$), ns, not significant.

NIL Growth Chamber Experiment

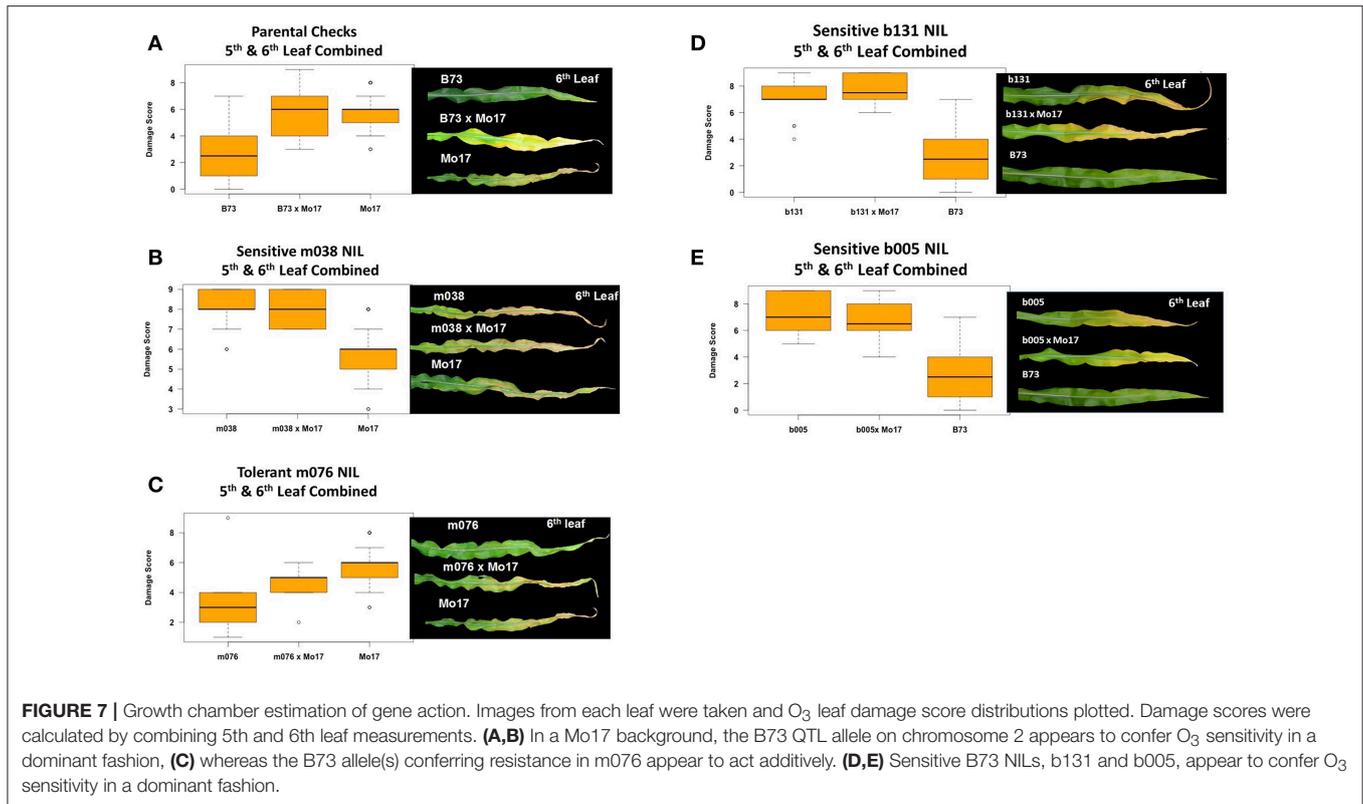
Experiments in the growth chambers with selected NILs and hybrids revealed that the leaf damage scores showed a binary distribution, and not the continuous degree of damage that was observed in the field (Figures 6A,C). Leaf damage scores on the fifth and sixth leaves were similar (Figures 6A,B). B73 and Mo17 were clearly distinguishable in both the field and the growth chamber. Mo17 NILs m002 and m016 carry the introgression chromosome 2. While, NIL m038 is heterozygous on chromosome 2 for the shared introgression under the QTL. Using leaf damage scores from NILs and hybrids in the chambers, we observed that the B73 QTL allele on chromosome 2 appeared to confer O₃ sensitivity in the Mo17 background in a dominant fashion (Figures 7A,B), whereas the B73 allele(s) conferring resistance in m076 appeared to act additively (Figure 7C). Sensitive B73 NILs, b005, and b131, appeared to confer O₃ sensitivity in a dominant fashion (Figures 7D,E). All of the sensitive NILs had a similar high leaf damage score (~8) in the growth chamber, despite showing a range of scores in the field.

DISCUSSION

This study used FACE technology to screen maize B73—Mo17 NILs in ambient and elevated [O₃] in order to identify genetic regions associated with O₃ response. Two NIL populations were screened, one with Mo17 as the recurrent background, and one with B73. Mo17 was more sensitive to elevated [O₃] as evidenced by greater visual leaf damage, and the NILs showed transgressive segregation (Figure 2). A significant QTL on chromosome 2 was identified and was consistent in both years of the study (Figure 4 and Table 1). The direction of the QTL effect was unexpected since B73 showed less leaf damage in elevated [O₃] than Mo17 (Figure 2). Just because B73 is more tolerant overall does not mean that it will have a tolerant allele for all QTL. Still, given the complexity of plant responses to elevated



[O₃], it is unexpected that only one major QTL was detected and that B73 introgressions into Mo17 make those NILs more susceptible. Other O₃ studies of Arabidopsis and rice have reported multiple smaller QTL for O₃ leaf damage (Shi et al., 2009; Brosché et al., 2010; Frei, 2015; Burton et al., 2016), and in rice, an apoplastic protein closely related to ascorbate oxidase, but lacking enzyme activity was subsequently suspected of involvement in O₃ tolerance (Ueda et al., 2015). There are several possible explanations for the maize results, including: (i) genetic background matters, (ii) there is cytoplasmic inheritance, or (iii) the B73 alleles have been unmasked. At this time it is inconclusive if the Mo17 introgression into B73 in this region has an effect. All NILs were derived from a B73 × Mo17 hybrid with B73 cytoplasm. An experiment using F₂ populations to determine whether the cytoplasm (B73 vs. Mo17) has an effect on the detection of the QTL effect could be revealing. F₂ populations could determine whether the QTL effect is only present in certain cytoplasm. To test for genetic background effects B73 NILs with a Mo17 introgression in this location could be leveraged. When particular natural variants are placed into different backgrounds the phenotypic consequences of that allele may be profoundly different from in their own background (Chandler et al., 2013). Genetic background effects have been



observed in most genetically tractable organisms where isogenic lines are used, including mice (Strunk et al., 2004), nematodes (Remold and Lenski, 2004), fruit flies (Gibson and van Helden, 1997), yeast (Dowell et al., 2010), rice (Cao et al., 2007), Arabidopsis (Huang et al., 2012), and bacteria (Wang et al., 2013).

Mapping is completed in general stages (Mackay et al., 2009). We identified a QTL that defines a large genomic region where one or more alleles affecting the trait segregate. In this study, the QTL associated with O₃ leaf damage on chromosome 2 has support intervals that crossed the centromere. This can reduce the probability of recovering recombinants in subsequent generations (Rodgers-Melnick et al., 2015) and make fine mapping more challenging. Still, theoretically, the next steps in fine mapping would be screen backcrossed F₂'s to identify segregating NILs. This would assist determination of which of the five introgressions is responsible for the O₃-induced leaf damage QTL. The shortcoming of such a classical QTL study is that the resolution of mapping is limited by the number of genetic recombination events occurring in the mapping populations (Lipka et al., 2015).

A second aim of this study was to test if growth chamber experiments could be used to accelerate screening for O₃ responsive lines. FACE technology, developed to study elevated CO₂ concentrations, is readily adaptable to study O₃. In fact, this has been used to study the impact of O₃ on various crops (reviewed in Ainsworth, 2017). There are certain limitations to all field trials including those using FACE technology. In many crop growing regions, only a single field season is possible, and FACE “rings” have a limited size. Given the desire for breeding programs to screen potentially thousands

of genotypes, conducting mapping experiments in smaller scale chamber experiments followed by verification in the field with elevated [O₃] may be more feasible (Frei, 2015). Our results showed that exposure to elevated O₃ in chambers caused leaf damage (Figure 5), and that O₃ sensitive and tolerant lines were consistent with field-based observations (Figure 6). In the growth chambers, damage was either present or absent, whereas in the field, we observed a continuum of leaf damage (Figure 6). This may be because we used a higher concentration of O₃ in the growth chambers compared to the field or because the air flow in the growth chambers provides greater O₃ exposure to the plants. Or alternatively that, the FACE experiment could capture small differences that could not be observed in chamber experiment. It was encouraging that the chamber experiments, even with differences in the degree of leaf damage, supported the field experiments overall.

We further used the growth chambers to examine potential dominance in sensitive and tolerant NILs (Figure 7). Mapping populations of Arabidopsis, poplar, rice and soybean have been used previously to identify QTL associated with O₃ response (Kim et al., 2004; Frei et al., 2008, 2010; Brosché et al., 2010; Street et al., 2011; Tsukahara et al., 2013, 2015; Burton et al., 2016). Brosché et al. (2010) identified three QTL on chromosomes 1, 2, and 3 for O₃ induced leaf injury in Arabidopsis. This research found that genes under these QTL are involved in ROS signaling and stomatal regulation. Street et al. (2011) screened a F₂ *Populus trichocarpa* × *Populus deltoids* population in a chronic O₃ study. They found QTL for leaf necrosis, diameter, late-season leaf number, height, late season abscission, area of the first unfurled leaf, and chlorophyll content in elevated [O₃]

or as response QTL. These findings indicate that O₃ responsive genomic regions exist and govern numerous traits. In soybean, Burton et al. (2016) found a leaf injury QTL following acute O₃ treatment varied with leaf position in the canopy. The analysis showed that loci were associated with distinct leaf developmental stages while the O₃ sensitive parent contributed one favorable allele for O₃ response. In rice O₃ damage QTL have been found on chromosomes 1, 2, 5, 7, 8, 9, and 11 and they tend to co-localize. Two rice QTL for leaf bronzing were further studied and found to contribute to O₃ tolerance via difference mechanisms (Frei et al., 2008, 2010; Shi et al., 2009). Breeding lines containing both bronzing QTLs were developed, and the lines were more O₃ tolerant than the sensitive line and the parental chromosome segment substitution lines.

To advance the findings reported here for maize, F₂ populations and markers have been developed. These resources can be utilized to further fine map the leaf damage QTL identified on chromosome 2 at 161 Mb. Additionally, they can be used to map sensitive B73 NILs, b007, and b003, and tolerant Mo17 NIL m076 down to a single introgression. Co-dominant markers can be used to screen for recombinants. These populations have been created to fine map resistance in m076, sensitivity in b005 and b131, and sensitivity in Mo17 NILs (m007, m022, m030, m072, and m091). These developed resources could be used to study which regions and/or genes are underpinning O₃ response in maize.

CONCLUSIONS

Ozone pollution is hypothesized to cost up to 10% of current maize yields in the U.S. (McGrath et al., 2015). Progress in

improving maize tolerance to oxidative stress requires genetic variability, reliable selection methods, time, and resources. Here, we identified a repeatable O₃-induced leaf damage QTL on chromosome 2 at 161Mb, which appears to confer O₃ sensitivity in a dominant manner. We further showed that O₃ sensitive lines identified in the field were also sensitive to elevated [O₃] in growth chambers, thus opening potential for high-throughput phenotyping and fine mapping of early season O₃ damage QTL in a controlled environment.

DATA AVAILABILITY

The raw data supporting the conclusions of this manuscript will be made available by the authors, without undue reservation, to any qualified researcher.

AUTHOR CONTRIBUTIONS

All authors listed have made a substantial, direct and intellectual contribution to the work, and approved it for publication.

FUNDING

This work was supported by the National Science Foundation under grant number PGR-1238030.

SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fsufs.2019.00051/full#supplementary-material>

REFERENCES

- Ainsworth, E. A. (2017). Understanding and improving global crop response to ozone pollution. *Plant J.* 90, 886–897. doi: 10.1111/tpj.13298
- Ainsworth, E. A., Serbin, S. P., Skoneczka, J. A., and Townsend, P. A. (2014). Using leaf optical properties to detect ozone effects on foliar biochemistry. *Photosynth. Res.* 119, 65–76. doi: 10.1007/s11120-013-9837-y
- Ainsworth, E. A., Yendrek, C. R., Sitch, S., Collins, W. J., and Emberson, L.D. (2012). The effects of tropospheric ozone on net primary productivity and implications for climate change. *Annu. Rev. Plant Biol.* 61, 637–661. doi: 10.1146/annurev-arplant-042110-103829
- Avnery, S., Mauzerall, D. L., Liu, J., and Horowitz, L. W. (2011). Global crop yield reductions due to surface ozone exposure: 1. Year 2000 crop production losses and economic damage. *Atmos. Environ.* 45, 2284–2296. doi: 10.1016/j.atmosenv.2010.11.045
- Bates, D., Maechler, M., Bolker, B., and Walker, S. (2015). Fitting linear mixed-effects models using lme4. *J. Stat. Softw.* 67, 1–48. doi: 10.18637/jss.v067.i01
- Betzberger, A. M., Gillespie, K. M., McGrath, J. M., Koester, R. P., Nelson, R. L., and Ainsworth, E. A. (2010). Effects of chronic elevated ozone concentration on antioxidant capacity, photosynthesis and seed yield of 10 soybean cultivars. *Plant Cell Environ.* 33, 1569–1581. doi: 10.1111/j.1365-3040.2010.02165.x
- Black, V. J., Black, C. R., Roberts, J. A., and Stewart, C. A. (2000). Tansley review no. 115, impact of ozone on the reproductive development of plants. *New Phytol.* 147, 421–447. doi: 10.1046/j.1469-8137.2000.00721.x
- Brace, S., Peterson, D. L., and Bowers, D. (1999). *A Guide to Ozone Injury in Vascular Plants of the Pacific Northwest*. US Department of Agriculture, Forest Service, Pacific Northwest Research Station.
- Brosché, M., Merilo, E., Mayer, F., Pechter, P., Puzörjova, I., Brader, G., et al. (2010). Natural variation in ozone sensitivity among *Arabidopsis thaliana* accessions and its relation to stomatal conductance. *Plant Cell Environ.* 33, 914–925. doi: 10.1111/j.1365-3040.2010.02116.x
- Burton, A. L., Burkey, K. O., Carter, T. E., Orf, J., and Cregan, P. B. (2016). Phenotypic variation and identification of quantitative trait loci for ozone tolerance in a Fiskeby III × Mandarin (Ottawa) soybean population. *Theor. Appl. Genet.* 129, 1113–1125. doi: 10.1007/s00122-016-2687-1
- Cao, Y., Ding, X., Cai, M., Zhao, J., Lin, Y., Li, X., et al. (2007). The expression pattern of a rice disease resistance gene Xa3/Xa26 is differentially regulated by the genetic backgrounds and developmental stages that influence its function. *Genetics* 177, 523–533. doi: 10.1534/genetics.107.075176
- Chandler, C. H., Chari, S., and Dworkin, I. (2013). Does your gene need a background check? How genetic background impacts the analysis of mutations, genes, and evolution. *Trends Genet.* 29, 358–366. doi: 10.1016/j.tig.2013.01.009
- Dowell, R. D., Ryan, O., Jansen, A., Cheung, D., Agarwala, S., Danford, T., et al. (2010). Genotype to phenotype: a complex problem. *Science* 328, 469–469. doi: 10.1126/science.1189015
- Eichten, S. R., Foerster, J. M., de Leon, N., Kai, Y., Yeh, C. T., Liu, S., et al. (2011). B73-Mo17 near-isogenic lines demonstrate dispersed structural variation in maize. *Plant Physiol.* 156, 1679–1690. doi: 10.1104/pp.111.174748
- FAO (2018). FAOSTAT. Available online at: <http://www.fao.org/faostat/en/#home> (accessed February 2018).
- Frei M., Tanaka J. P., Chen C., and Wissuwa M. (2010). Mechanisms of ozone tolerance in rice: characterization of two QTLs affecting leaf bronzing by gene expression profiling and biochemical analyses. *J. Exp. Bot.* 61, 1405–1417.
- Frei, M. (2015). Breeding of ozone resistant rice: relevance, approaches and challenges. *Environ. Pollut.* 197, 144–155. doi: 10.1016/j.envpol.2014.12.011

- Frei, M., Tanaka, J. P., and Wissuwa, M. (2008). Genotypic variation in tolerance to elevated ozone in rice: dissection of distinct genetic factors linked to tolerance mechanisms. *J. Exp. Bot.* 59, 3741–3752. doi: 10.1093/jxb/ern222
- Gibson, G., and van Helden, S. (1997). Is function of the Drosophila homeotic gene Ultrathorax canalized? *Genetics* 147, 1155–1168.
- Huang, X., Effen, S., Meyer, R. C., Theres, K., and Koornneef, M. (2012). Epistatic natural allelic variation reveals a function of AGAMOUS-LIKE6 in axillary bud formation in Arabidopsis. *Plant Cell* 24, 2364–2379. doi: 10.1105/tpc.112.099168
- Kangasjarvi, J., Jaspers, P., and Kollist, H. (2005). Signaling and cell death in ozone-exposed plants. *Plant Cell Environ.* 28, 1021–1036. doi: 10.1111/j.1365-3040.2005.01325.x
- Kay, D. K., Mueller, N. D., West, P. C., and Foley, J. A. (2013). Yield trends are insufficient to double global crop production by 2050. *PLoS ONE* 8:e66428. doi: 10.1371/journal.pone.0066428
- Kim, K. M., Kwon, Y. S., Lee, J. J., Eun, M. Y., and Sohn, J. K. (2004). QTL mapping and molecular marker analysis for the resistance of rice to ozone. *Mol. Cells* 17, 151–155.
- Krupa, S., McGrath, M. T., Andersen, C. P., Booker, F. L., Burkey, K. O., Chappelka, A. H., et al. (2001). Ambient ozone and plant health. *Plant Dis.* 85, 4–12. doi: 10.1094/PDIS.2001.85.1.4
- Leisner, C. P., and Ainsworth, E. A. (2012). Quantifying the effects of ozone on plant reproductive growth and development. *Glob. Chang. Biol.* 18, 606–616. doi: 10.1111/j.1365-2486.2011.02535.x
- Lipka, A. E., Kandianis, C. B., Hudson, M. E., Yu, J., Drnevich, J., Bradbury, P. J., et al. (2015). From association to prediction: statistical methods for the dissection and selection of complex traits in plants. *Curr. Opin. Plant Biol.* 24, 110–118. doi: 10.1016/j.pbi.2015.02.010
- Mackay, T. F., Stone, E. A., and Ayroles, J. F. (2009). The genetics of quantitative traits: challenges and prospects. *Nat. Rev. Genet.* 10, 565–572. doi: 10.1038/nrg2612
- McGrath, J. M., Betzelberger, A. M., Wang, S., Shook, E., Zhu, X. G., Long, S. P., et al. (2015). An analysis of ozone damage to historical maize and soybean yields in the United States. *Proc. Natl. Acad. Sci. U.S.A.* 112, 14390–14395. doi: 10.1073/pnas.1509777112
- Miller, J. E., Heagle, A. S., Vozzo, S. F., Philbeck, R. B., and Heck, W. W. (1989). Effects of ozone and water-stress, separately and in combination, on soybean yield. *J. Environ. Qual.* 18, 330–336. doi: 10.2134/jeq1989.00472425001800030016x
- Mills, G., Sharps, K., Simpson, D., Pleijel, H., Broberg, M., Uddling, J., et al. (2018). Ozone pollution will compromise efforts to increase global wheat production. *Glob. Chang. Biol.* 24, 3560–3574. doi: 10.1111/gcb.14157
- Morgan, P. B., Bernacchi, C. J., Ort, D. R., and Long, S. P. (2004). An *in vivo* analysis of the effect of season-long open-air elevation of ozone to anticipated 2050 levels on photosynthesis in soybean. *Plant Physiol.* 135, 2348–2357. doi: 10.1104/pp.104.043968
- Pachauri, R. K., Allen, M. R., Barros, V. R., Broome, J., Cramer, W., Christ, R., et al. (2014). *Climate Change 2014: Synthesis Report*. Contribution of Working Groups I, II and III to the fifth assessment report of the Intergovernmental Panel on Climate Change (IPCC), 151.
- Perez, I. B., and Brown, P. J. (2014). The role of ROS signaling in cross-tolerance: from model to crop. *Front. Plant Sci.* 5, 1–6. doi: 10.3389/fpls.2014.00754
- R Core Team (2015). *R: A Language and Environment for Statistical Computing*. Vienna: R Foundation for Statistical Computing.
- Rao, M. V., and Davis, K. R. (1999). Ozone-induced cell death occurs via two distinct mechanisms in Arabidopsis: the role of salicylic acid. *Plant J.* 17, 603–614. doi: 10.1046/j.1365-313X.1999.00400.x
- Rao, M. V., and Davis, K. R. (2001). The physiology of ozone induced cell death. *Planta* 213, 682–690.
- Rao, M. V., Lee, H. I., Creelman, R. A., Mullet, J. E., and Davis, K. R. (2000). Jasmonic acid signaling modulates ozone-induced hypersensitive cell death. *Plant Cell* 12, 1633–1646. doi: 10.1105/tpc.12.9.1633
- Remold, S. K., and Lenski, R. E. (2004). Pervasive joint influence of epistasis and plasticity on mutational effects in *Escherichia coli*. *Nat. Genet.* 36, 423–437. doi: 10.1038/ng1324
- Rodgers-Melnick, E., Bradbury, P. J., Elshire, R. J., Glaubitz, J. C., Acharya, C. B., Mitchell, S. E., et al. (2015). Recombination in diverse maize is stable, predictable, and associated with genetic load. *Proc. Natl. Acad. Sci. U.S.A.* 112, 3823–3828. doi: 10.1073/pnas.1413864112
- Shi, G., Yang, L., Wang, Y., Kobayashi, K., Zhu, J., Tang, H., et al. (2009). Impact of elevated ozone concentration on yield of four Chinese rice cultivars under fully open-air field conditions. *Agric. Ecosyst. Environ.* 131, 178–184. doi: 10.1016/j.agee.2009.01.009
- Street, N. R., James, T. M., James, T., Mikael, B., Jaakko, K., Mark, B., et al. (2011). The physiological, transcriptional and genetic responses of an ozone-sensitive and an ozone-tolerant poplar and selected extremes of their F2 progeny. *Environ. Pollut.* 159, 45–54. doi: 10.1016/j.envpol.2010.09.027
- Strunk, K. E., Amann, V., and Threadgill, D. W. (2004). Phenotypic variation resulting from a deficiency of epidermal growth factor receptor in mice is caused by extensive genetic heterogeneity that can be genetically and molecularly partitioned. *Genetics* 167, 1821–1832. doi: 10.1534/genetics.103.020495
- Tang, H., Liu, G., Han, Y., Zhu, J., and Kobayashi, K. (2011). A system for free-air ozone concentration elevation with rice and wheat: control performance and ozone exposure regime. *Atmos. Environ.* 45, 6276–6282. doi: 10.1016/j.atmosenv.2011.08.059
- Tsukahara, K., Sawada, H., Kohno, Y., Matsuura, T., Mori, I. C., Terao, T., et al. (2015). Ozone-induced rice grain yield loss is triggered via a change in panicle morphology that is controlled by ABERRANT PANICLE ORGANIZATION 1 gene. *PLoS ONE* 10:e0123308. doi: 10.1371/journal.pone.0123308
- Tsukahara, K., Sawada, H., Matsumura, H., Kohno, Y., and Tamaoki, M. (2013). Quantitative trait locus analyses of ozone-induced grain yield reduction in rice. *Environ. Exp. Bot.* 88, 100–106. doi: 10.1016/j.envexpbot.2011.12.012
- Ueda, Y., Siddique, S., and Frei, M. (2015). A novel gene, OZONE-RESPONSIVE APOPLASTIC PROTEIN1, enhances cell death in ozone stress in rice. *Plant Physiol.* 169, 873–889.
- Van Dingenen, R., Dentener, F. J., Raes, F., Krol, M. C., Emberson, L., and Cofala, J. (2009). The global impact of ozone on agricultural crop yields under current and future air quality legislation. *Atmos. Environ.* 43, 604–618. doi: 10.1016/j.atmosenv.2008.10.033
- Wang, Y., Arenas, C. D., Stoebel, D. M., and Cooper, T. F. (2013). Genetic background affects epistatic interactions between two beneficial mutations. *Biol. Lett.* 9:20120328. doi: 10.1098/rsbl.2012.0328
- Yendrek, C., Tomaz, T., Montes, C. M., Cao, Y., Morse, A. M., Brown, P. J., et al. (2017). High-throughput phenotyping of maize leaf physiology and biochemistry using hyperspectral reflectance. *Plant Physiol.* 173, 614–626. doi: 10.1104/pp.16.01447
- Yendrek, C. R., Erice, G., Montes, C. M., Tomaz, T., Sorgini, C. A., Brown, P. J., et al. (2017). Elevated ozone reduces photosynthetic carbon gain by accelerating leaf senescence of inbred and hybrid maize in a genotype-specific manner. *Plant Cell Environ.* 40, 3088–3100. doi: 10.1111/pce.13075
- Zhu, X., Feng, Z., Sun, T., Liu, X., Tang, H., Zhu, J., et al. (2011). Effects of elevated ozone concentration on yield of four Chinese cultivars of winter wheat under fully open-air field conditions. *Glob. Chang. Biol.* 17, 2697–2706. doi: 10.1111/j.1365-2486.2011.02400.x

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