Different effects of phytohormones on Fusarium head blight and Fusarium root rot resistance in *Brachypodium distachyon*

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The tissue-dependent effects of defence phytohormones and tissue-independent effects of development phytohormones on *F. graminearum*-induced Fusarium head blight and Fusarium root rot diseases in the model cereal *Brachypodium distachyon*.

Abstract

Fusarium graminearum is a devastating pathogen of small grain cereals causing both Fusarium head blight (FHB), Fusarium root rot (FRR). Phytohormones play important roles in hosts during Fusarium infection and studies have found that application of phytohormones can significantly impact FHB resistance. In contrast to FHB, FRR remains a poorly characterised disease and it is unknown whether phytohormones play similar roles in FHB and FRR. In this present study, *B. distachyon* floral tissues at mid anthesis and root tissues from seedlings were exogenously treated with a range of phytohormones before inoculation with *F. graminearum*. Application of salicylic acid (SA) significantly increased susceptibility to FRR but not to FHB. On the other hand, both jasmonic acid (JA) and the ethylene precursor 1-aminocyclopropane-1-carboxylic acid (ACC) increased resistance to FRR but increased susceptibility to FHB. The auxins indole-3-acetic acid (IAA) and 1-napthaleneacetic acid (NAA) greatly increased FHB and FRR resistance. The cytokinin trans-Zeatin and the defence-related compound 3-aminobutanoic acid (BABA) induced high levels of susceptibility in both FHB and FRR. In FRR the phytohormones SA, JA, and ACC appeared to exert their effects early in infection whereas auxin, cytokinins, and BABA altered the rate of colonisation rather than affecting initial infection.

Key Words

Abbreviations

ACC, 1-aminocyclopropane-1-carboxylic acid; BABA, 3-aminobutanoic acid; dpi, days post inoculation; FHB, Fusarium head blight; FRR, Fusarium root rot; IAA indole-3-acetic acid; JA, jasmonic acid; NAA, 1-napthaleneacetic acid; PDA, potato dextrose agar; RNL, root necrosis length; SA, salicylic acid.

Introduction

Fusarium graminearum is one of the most important plant pathogens in the world (Dean et al., 2012), affecting small grain cereals such as bread wheat (T. aestivum), barley (H. vulgare), and rye (S. cereale). F. graminearum can infect almost the entire plant (Miedaner, 1997) causing a range of diseases. The most well-known and economically important disease is fusarium head blight (FHB). Other diseases include fusarium crown rot, fusarium root rot (FRR), and seedling blight. Plant pathogens often adopt specific trophic lifestyles. Biotrophs obtain nutrients from living tissue, necrotrophs from dead tissue, and hemibiotrophic pathogens adopt an early biotrophic phase that is followed by a necrotrophic phase sometime later in infection (Glazebrook, 2005; Zeilinger et al., 2016). F. graminearum is generally considered a facultative hemibiotroph in the developing cereal spike (Jansen et al., 2005; Boddu et al., 2006). Infection patterns of floral tissues show that F. graminearum grows as a biotroph at the advancing hyphal front with necrotrophic feeding behind this, where hyphae are ramifying through tissues already killed by the fungus (Brown et al., 2010). Current FHB control methods rely on the effective use of fungicides and genetics resistance. However novel control mechanisms are required.

Fungal root diseases are widespread in wheat and barley fields and are becoming more prevalent with the increased use of cereal crop rotation and no tillage practices (Cook, 2001). FRR is caused by several Fusarium species including F. graminearum. Colonisation and sporulation can occur rapidly (Wang et al., 2015) causing root browning and necrosis (Beccari et al., 2011). FHB and Fusarium crown rot can also manifest from FRR infection due to systemic migration via the vascular system (Beccari et al., 2011; Wang et al., 2015). Ultimately this results in reduced root, shoot length, biomass and yield loss (Mergoum et al., 1998; Beccari et al., 2011; Wang et al., 2015). Control of FRR is also difficult due to limited effectiveness of fungicides, lack of genetic resistance, and persistence in the soil for many years facilitated by the predominantly saprotrophic lifestyle of F. graminearum (Cook, 2001). Furthermore, FRR is not as well characterised as FHB due to the difficulty of research on root diseases.

Brachypodium distachyon (purple false brome) is a valuable model for small-grained

cereals as it is a small inbreeding annual with short generation times, minimal growth

requirements, and abundant genetic resources (Draper *et al.*, 2001; Vogel *et al.*, 2006; Vogel

et al., 2010; Brkljacic *et al.*, 2011; Kellogg, 2015; Scholthof *et al.*, 2018). *B. distachyon* is a

good model for investigating shoot and root development due to high anatomical and

developmental similarities to wheat (Draper *et al.*, 2001; Watt *et al.*, 2009; Opanowicz *et al.*,

2011; Kellogg, 2015). Furthermore, hormone signalling pathways appear well conserved

(Goddard *et al.*, 2014; Kouzai *et al.*, 2016). Importantly, *B. distachyon* is an excellent model

for investigating FHB and FRR as both roots and florets are susceptible to *F. graminearum*

infection (Peraldi *et al.*, 2011; Pasquet *et al.*, 2014).

Phytohormones are critical components of plant defence signalling and their impact on *F.*

graminearum resistance response has been investigated in numerous studies on *A. thaliana*,

wheat, and barley. Exogenous application of salicylic acid (SA) or methylsalicylate has been

shown to have positive effects on FHB resistance in wheat (Makandar *et al.*, 2011; Qi *et al.*,

2012; Sorahinobar *et al.*, 2016) with direct effects on *in-vitro* growth (Qi *et al.*, 2012).

Application of methylsalicylate also showed positive effects on wheat leaf resistance to *F.*

graminearum (Ameye *et al.*, 2015). Jasmonic acid (JA) and methyl-jasmonic acid application

was also reported to have a positive effect on FHB resistance in wheat (Li and Yen, 2008; Qi

et al., 2016; Sun *et al.*, 2016). In contrast Makandar and colleagues (2010) found that

methyl-jasmonic acid compromised resistance during early infection but had positive effects

during late infection in *A. thaliana* (Makandar *et al.*, 2010). Similar results were reported in

wheat where pre-infection application of methyl-jasmonic acid promoted susceptibility,

whereas post-infection application promoted resistance of wheat leaves to *F. graminearum*

(Ameye *et al.*, 2015). JA was also found to reduce *F. graminearum* growth in vitro (Qi *et al.*,

2016). The precise role of ethylene on resistance to Fusarium remains unclear. Compounds

that produce ethylene have been reported to have positive effects (Li and Yen, 2008; Foroud

et al., 2018), negative effects (Chen *et al.*, 2009), and no significant effect (Sun *et al.*, 2016)

on Fusarium infection.

More recently other important hormones were shown to have roles in defence and

contribute significantly to the infection response. Exogenous application of gibberellic acid

reduced FHB spread in wheat heads (Buhrow *et al.*, 2016), whereas exogenous application

of abscisic acid negatively affected resistance to FHB (Buhrow *et al.*, 2016; Qi *et al.*, 2016).

Exogenous application of brassinosteroid had positive effects on resistance (Ali *et al.*, 2013)

while brassinosteroid receptor mutants in which brassinosteroid levels are believed to be

increased, exhibited enhanced resistance to *Fusarium culmorum* (Goddard *et al.*, 2014).

Finally, the auxin indole-3-acetic acid (IAA) had positive effects on barley FHB (Petti *et al.*,

2012), and was even found to reduce Fusarium growth *in vitro* (Luo *et al.*, 2016). There are

hormones that have not been investigated regarding exogenous application and response to

F. graminearum infection. These include cytokinins which have been implicated in plant

defence (Choi *et al.*, 2010; Choi *et al.*, 2011; Albrecht and Argueso, 2017) and the recently

classed non-protein amino acid and signalling molecule 3-aminobutryric acid (BABA) (Cohen,

2001; Jakab *et al.*, 2001; Cohen, 2002; Cohen *et al.*, 2016; Thevenet *et al.*, 2017).

The reported effects of phytohormones on resistance to *F. graminearum* is complex or in

some cases contradictory. Most studies have focused on FHB with a lack of research on *F.*

graminearum-induced FRR pathogenesis. The main aim of this research was to investigate

the effect of several phytohormones on resistance to FRR and compare these with the

effects of the same compounds on FHB. We then generated a model for how each

phytohormone influences resistance or susceptibility to the two diseases in *B. distachyon*.

Materials and Methods

Plant Material and Growth Conditions.

The *Brachypodium distachyon* line Bd3-1 was obtained from the John Innes Centre, Norwich, UK. To soften the floret, seeds were soaked in water for 10-30 min. Subsequently, the lemma and palea were peeled off the individual seeds. Seeds were placed between two layers of dampened filter paper (9 cm, Sartorius Grade 292) with 5 ml sterile water, and were stratified at 5 °C for five days in the dark, and incubated for one day at 22 °C (16h/8h light/dark photoperiod, variable humidity) in controlled environment growth cabinets (Peraldi *et al.*, 2011). For FHB assays, peeled Bd3-1 seeds were sown in 50% Peat/Sand and 50% John Innes mix 2 (two seeds per pot). Plants were kept for four weeks at 22°C (20h/4h light/dark photoperiod, 70% humidity) in controlled environment growth cabinets or rooms (Peraldi *et al.*, 2011). The *B. distachyon* line Bd21 was obtained from BASF SE Agricultural Centre Limburgerhof, Germany, For the FHB assay, approximately five Bd21 unpeeled seeds were grown for 6 weeks at 20h/4h (day/night) in a glasshouse (Autumn conditions).

Preparation and Maintenance of *Fusarium graminearum*

The *Fusarium graminearum* isolate PH1 was used for all experiments. In order to produce mycelial inoculum for FRR assays, *F. graminearum* was maintained on approximately 20 ml potato dextrose agar (PDA) in 9 cm diameter plastic Petri-dishes in a controlled growth cabinet at 22°C under 16h/8h light/dark photoperiod. Conidial *F. graminearum* inoculum was produced in Mung Bean broth as reported previously (Makandar *et al.*, 2006). After one week, conidia were harvested, washed and concentration adjusted to 1 x 10^6 conidia/ml. Inoculum was stored at 5 °C for up to two weeks before inoculation. For Bd21 FHB assay, *F. graminearum* isolate Li600 was grown alternating on malt agar and oatmeal agar for two weeks and conidia were harvested in 10 ml of water amended with 0.05 % Tween 20 and concentration adjusted as above.

Fusarium Root Rot Assays with Chemical Amendment.

Aspects of the FRR assay were derived from (Peraldi *et al.*, 2011; Goddard *et al.*, 2014)

and modified for chemical amendment experiments. A sterile 9 cm^2 filter paper square (cut

from Chromatography Paper 46 cm x 57 cm from Slaughter Ltd R & L) was placed on square

plastic square Petri-dishes (10 cm^2) containing 50 ml autoclaved 0.8% agar (Fischer Science).

Under sterile conditions, ten cold-stratified and germinated Bd3-1 seeds were placed on the

filter paper. A minimum of 30 seedlings was used for each treatment. All plates were placed,

angled at 70° from the horizontal to ensure uniform downward root growth, in covered

plant propagators containing wetted paper towel to maintain high humidity (Supp Mat. Fig.

x). Plant propagators with square Petri-dishes were incubated at 22 °C (16h/8h light/dark

photoperiod, variable humidity) in controlled environment growth cabinets. After three

days, the filter paper with seedlings attached was carefully transferred to different square

Petri-dishes containing 0.8% agar amended with phytohormone or control solvent alone

and returned to controlled environment growth cabinets. All compounds were ordered

from Merck/Sigma-Aldrich UK unless otherwise stated. The final concentration for SA, JA,

ACC, and 3-aminobutanoic acid (BABA) were derived from a dilution series experiment

including (Kakei *et al.*, 2015) and (Peraldi, unpublished). The concentration for IAA was

derived from (Kakei *et al.*, 2015) whereas trans-Zeatin was derived from (Großkinsky *et al.*,

2013). The concentrations for the 1-napthaleneacetic acid (NAA) and kinetin were derived

from the concentrations chosen from IAA and trans-Zeatin respectively. The compounds

ACC, IAA – sodium salt, kinetin, and BABA were dissolved in water. The compounds SA, JA

and NAA were dissolved in ethanol and trans-zeatin was dissolved in DMSO. All non-water

solvent concentrations in final treatment were at or below 0.1 %. The same concentration of

solvent was applied to respective control treatment 0.8 % agar plates.

Roots were inoculated six hours after transfer to hormone amended medium. Inoculum

was prepared by blending mycelium and PDA from one-week old cultures (adding 1 ml

water per PDA plate). Subsequently, approximately 0.1 ml to 0.2 ml of homogenized slurry

was transferred with a 10 ml syringe onto the root tip. This mycelial slurry induced more

severe and consistent symptoms than a conidial slurry (PDA slurry amended with a conidial

suspension) during early FRR (Unpublished data). Inoculated plants were then incubated at

22 °C (16h/8h light/dark photoperiod, variable humidity) in controlled growth cabinets. The

slurry was removed when necrosis was visible at the root tip, often at 1-day post inoculation

(dpi). Roots were photographed at intervals to monitor disease development. For repeated

independent experiments with different measurement dates, the days were combined and

denoted as 'Score Date' 1 to 3 with each measurement at 3-day intervals (Fig. 1B, Fig. 1C).

Fusarium Head Blight Assays with Chemical Pre-Treatment.

193   Numerous features of the FHB assay were derived from (Peraldi *et al.*, 2011). With

Bd3-1 FHB assays, once extruding anthers were visible around mid-anthesis, the entire Bd3-

1 plant was sprayed with 50 ml of phytohormone or solvent with 0.05% Tween 20 onto a

tray of 10 to 11 pots. All compounds were ordered from Merck/Sigma-Aldrich UK unless

otherwise stated. The concentrations used for BABA and trans-Zeatin were derived from

those used in FRR assays, the SA concentrations was obtained from previous publications

(Mandal *et al.*, 2009; Makandar *et al.*, 2011; Sorahinobar *et al.*, 2016) and JA and ACC were

experimentally determined. ACC and BABA were dissolved in water, trans-Zeatin in DMSO,

and SA and JA in ethanol. The final concentration of non-water solvents was kept at or

below 0.2 %. The same concentration of solvent was applied to respective control treatment

groups. Twenty-four hours later the soil was watered, and Bd3-1 spikes were sprayed to

run-off with conidia of *F. graminearum* ($0.25\text{-}1 \times 10^6$) amended with wetting agent (0.05%

Tween 20). A control pot without inoculum was included for each treatment. Approximately

30 ml of inoculum was sprayed onto a total of 20 plants' spikelets per treatment. Spraying

was performed immediately before the dark period and pots were then collectively held in a

large plastic humidity chamber to maximise humidity for three days 22°C (20h/4h light/dark

photoperiod). Symptoms were scored every four days by counting the number of infected

florets per spike. For repeated independent experiments with different measurement dates,

the days were combined to 'Score Date' 1-3 with each measurement at 4-day intervals (Fig.

3B, Fig. 3D).

Some experiments were performed at BASF SE using Bd21 instead of Bd3-1. Bd21 and

Bd3-1 are both highly susceptible to FHB (Peraldi *et al.*, 2011). For the Bd21 auxin trial, the

entire plant was sprayed with respective compounds until run-off. The auxin concentration

was experimentally determined (Unpublished data). The final solvent concentration for each

auxin treatment with Bd21 including the control treatment was 5 % acetone. After 24h, a

working concentration of 5×10^5 *F. graminearum* LI600 was prepared in 0.05 % tween 20

water, and 175 ml of inoculum was evenly sprayed above all plants. The base matting was

watered until run-off and plants were encased in plastic covers for six days in an elevated

humidity glasshouse room (Autumn conditions) before scoring.

<u>Statistical Analysis</u>.

All statistical tests were performed using the software package GENSTAT

v.19.1.0.21390 (VSN international Ltd). A Generalised liner model with an accumulated

analysis of variance (ANOVA) was used for all experiments using a normal distribution for

FRR assays or log-linear modelling for FHB assay count data. Individual time points were

analysed separately. All data was untransformed except for a square root transformation for

Fig. 1A at 7 dpi, Fig. 1C at score date 1, Fig. 2A at 2 dpi, Fig. 2C at 4 dpi, Fig. 2D at 2dpi and

4dpi. A log10 transformation was used for Fig. 2C at 2dpi. A few ANOVA's for combined

independent experiments showed a significant interaction between the independent

experiment and the hormone treatment at specific time points: Fig. 1A at 3 dpi, Fig. 1B at

Score Date 3, Fig. 1D at 2 dpi, Fig. 2A at 2 dpi, Fig. 2B at 2 dpi, Fig. 3C at 11 dpi, and Fig. 3E at

7 dpi and 11dpi ($p < .05$). However for all the remaining time points, no such interaction was

observed ($p > .05$). For Bd21, a randomised complete block design was used from the R

package 'agricolae' and plant pots were randomized before chemical application. Microsoft

Office was used for writing, data collection, diagrams, and analysis. All graphs were

prepared using Graphpad Prism (V5.04).

Results

The Differential Effect of Plant Hormones on Fusarium Root Rot Disease Progression.

The SA amendment significantly increased root necrosis length (RNL) at all time points (Fig. 1A), while amendment with JA (Fig. 1B), or the ethylene precursor (ACC) (Fig. 1C), significantly reduced RNL at all time points (p < .01). Interestingly, the difference in RNL for all three compounds remained similar with the regression lines remaining parallel rather than diverging over time. The compound BABA had the greatest effect in increasing RNL compared to other defence-related hormones (Fig. 1D). The differential RNL symptoms resulting from pre-application of BABA increased over time with application leading to near doubling of necrosis at 4 dpi and 6 dpi (p < .001) relative to the control. Overall exogenous application of defence-related hormones had a substantial effect on FRR disease severity but whether the effect on resistance was positive or negative was dependant on the particular phytohormone.

Exogenous application of IAA resulted in the most pronounced decrease in RNL compared to the control at all time points (p < .001) since FRR symptoms were reduced to approximately half those in control treatments (Fig 2A). Like IAA, the synthetic auxin NAA also reduced RNL at all time points (p < .001, Fig 2B) but auxin induced FRR resistance was most pronounced with IAA. With both auxins, the increased resistance was observed as early as 2 dpi yet the regression lines diverged over time with a greater reduction in RNL at later time points (Supp. Fig. S1). Overall, auxin treatment produced a very positive effect on resistance towards FRR.

The cytokinin trans-Zeatin increased RNL at all time points (p < .001, Fig. 2C) and the differential increased over time. The extent of necrosis was twice that in the control by 6 dpi (Supp. Fig. S2) leading to the most severe symptoms observed with any of the phytohormone treatments. The cytokinin kinetin exhibited no significant effect on symptoms at 2dpi (p = .785) but increased the rate of RNL so that the differential increased over time ((4 dpi and 6 dpi p < .01, p < .001 respectively) Fig. 2D). Strikingly the effect of cytokinin amendment increased dramatically over time, with the effect being even greater than that observed for the positive effect of auxins. Overall cytokinin amendment appeared

to have a very negative effect on FRR resistance which became more pronounced as the
disease progressed.

The phytohormones abscisic acid, gibberellic acid, and the brassinosteroid epibrassinolide
were also tested on FRR (Supp. Fig. S3A and S3B and Supp. Fig. S4). Gibberellic acid and
epibrassinolide had no significant effect on FRR symptoms while abscisic acid induced
extensive root discolouration that was indistinguishable from FRR symptoms and so
prevented assessment (Supp. Fig. S4).

Most Plant Hormones Promoted Susceptibility to Fusarium Head Blight.

In order to compare the effects of hormones on FRR to those on FHB, the same
hormones were applied to the entire *B. distachyon* plant and subsequently the spikes were
inoculated with *F. graminearum* conidia. Unless otherwise stated, the effects described are
from a single dose of phytohormone pre-treatment as opposed to continual exposure as
during the FRR assays. As a result, often a higher concentration of phytohormone was
applied to maximise the dose of phytohormone received by the plant.

Although pre-inoculation treatment with SA marginally increased FHB symptoms (Fig.
3A), the increase was not statically significant at any time point ($p > .05$). In additional
experiments, SA was applied repeatedly (four applications) before, during, and after
inoculation on Bd3-1 but again, no significant effect on resistance was observed (Supp. Fig.
S5). Pre-inoculation treatment with JA resulted in an increase in the number of infected
florets at all time points ($p < .05$, Fig. 3B). Similarly, pre-treatment of ACC (Fig. 3C) also
significantly increased the number of infected florets at all time points ($p < .05$), but the
effect on symptoms diminished over time and by 11 dpi, the effect was not significant ($p >$
.05). Pre-inoculation application of BABA significantly increased the number of infected
florets at all time points ($p < .001$) (Fig. 3D) and application of cytokinin trans-Zeatin caused
the largest increase in infected florets relative to the control treatment of any of the
compounds examined (Fig. 3E) ($p < .001$). With both BABA and trans-Zeatin, the differential
in the number of infected florets relative to the control treatments increased after the first
score date although the divergence from the controls was not as great as observed in the
FRR assays where the phytohormone amendment was present throughout the course of the
experiment. Overall pre-inoculation application of all the phytohormones except for SA

promoted FHB susceptibility leading to a significant increase in the number of florets
exhibiting symptoms.

Using Bd21 instead of Bd3-1, two concentrations of IAA and NAA were exogenously
applied to Bd21 plants before inoculation with the isolate *F. graminearum* LI6000 (Fig. 4). All
auxin pre-inoculation applications resulted in a reduction in the number of infected florets.
Pre-inoculation application of synthetic auxin NAA at 8 mM resulted in the most significant
decrease in the number of infected florets ($p < .001$) and even 4 mM NAA significantly
decreased the number of infected florets ($p < .01$). IAA appeared to be less effective than
NAA since 8 mM IAA showed a similar reduction in infected florets as 4 mM NAA ($p < .01$)
and at 4 mM IAA the reduction in the number of infected florets was not significant ($p =$
$.33$). Overall, these data indicate that auxins have a positive effect on FHB resistance when
applied at relatively high concentrations prior to inoculation.

To ascertain whether the effects of phytohormones on FRR reflected an altered plant
response or an effect on the fungus itself, hormones at the same concentrations as used in
the experiments were applied to *F. graminearum* growth medium to measure any changes
in growth over time. Compared to the respective control treatments there was no
significant effect on *F. graminearum* growth of any of the tested phytohormone (Supp. Fig.
S6). Due to the high concentrations of auxins tested for FHB, a different assay was
performed. Auxins or solvent controls were applied to filter disks at four equidistant points
from a mycelial plug of *F. graminearum* PH1 growing on PDA. There was no difference in
mycelial growth near any of the auxin-treated filter disks (Supp. Fig. S7).

Discussion

Given some of the proposed differences between *F. graminearum* infection of floral and root tissues, the effect of a range of phytohormones on FRR was assessed and compared to their effect on FHB. Research into the role of exogenous phytohormones on FHB resistance has been investigated in several plant species. SA was found to increase resistance to FHB in wheat and Arabidopsis (Makandar *et al.*, 2011; Qi *et al.*, 2012; Sorahinobar *et al.*, 2016). We were not able to replicate the positive FHB effect of SA (Fig. 3A) despite using similar concentrations to those used previously. We speculate that the different results are due to alternative SA application methods. In earlier studies, SA (200 µM) was applied as a soil drench (Makandar *et al.*, 2011) (Sorahinobar *et al.*, 2016). This application method may have more pronounced effects on resistance due to initial uptake in the roots as opposed to florets and because the phytohormone remained in contact with the roots throughout the experiment. Furthermore prolonged exposure to SA may influence JA/ethylene dependent Induced systemic resistance (ISR) (Pieterse *et al.*, 2014) which might impact the resistance response to FHB.

Our results on the effect of application of JA on FHB (Fig. 3B) support some studies (Makandar *et al.*, 2010), but contradict others (Li and Yen, 2008; Qi *et al.*, 2016; Sun *et al.*, 2016). The reasons for these differences are unclear. They may reflect the use of a higher concentration of JA in the present study than was used in some previous reports (Qi *et al.*, 2016; Sun *et al.*, 2016) or they may be due to differences in the timing of treatment (Makandar *et al.*, 2010). The difference may also be due to the application method. Qi and colleagues (2016) point-inoculated individual florets with *F. graminearum* rather than spraying the head (Qi *et al.*, 2016). Injecting high levels of inoculum into florets may have resulted in a by-passing or severe shortening of the biotrophic phase of infection/colonisation (Jansen *et al.*, 2005; Brown *et al.*, 2010). Similar to JA, our results with ACC (Fig. 3C) support the findings from some studies (Chen *et al.*, 2009), but not those of others (Li and Yen, 2008; Sun *et al.*, 2016; Foroud *et al.*, 2018). Ethylene can affect host resistance via synergism with classic defence hormones (Pieterse *et al.*, 2012) or via accelerated senescence (Abeles *et al.*, 2012; Häffner *et al.*, 2015). As a result, the outcome is very sensitive to several factors such as timing of exogenous application, tissue type, and

the type of invading pathogen (van Loon *et al.*, 2006). Differences in the experimental

procedures may also affect important factors for ethylene response. In some reports

detached head assays were performed (Chen *et al.*, 2009; Foroud *et al.*, 2018) which might

influence senescence processes as well as defence responses. Ethephon (releases ethylene

gas) was used instead of ACC as the ethylene treatment in some studies (Li and Yen, 2008;

Chen *et al.*, 2009; Sun *et al.*, 2016). Although Foroud and colleagues (2018) found that

ethephon and ACC had similar effects on FHB in most wheat varieties tested (Foroud *et al.*,

2018). The role of ethylene is further complicated given evidence that some isolates of *F.*

graminearum and *F. culmorum* can produce ethylene *in vitro* (Haidoulis et al. unpublished

data), however it is uncertain whether ethylene is produced *in planta*.

We found that auxin, (both IAA and NAA) greatly improved FHB resistance (Fig. 4). This

results supports evidence by Petti and colleagues (2012) who found that exogenous IAA

application reduced yield loss and general symptoms of *F. culmorum*-induced FHB in barley

(Petti *et al.*, 2012). However, the mechanism for the effects of auxin in FHB response are not

clear given that *F. graminearum* can also produce IAA (Luo *et al.*, 2016).

Overall, our results using *B. distachyon* as a model for investigating FHB responses were

broadly in line with those from studies on wheat, barley and Arabidopsis. Similar trends

were observed in the response to auxins, JA, and ethylene as reported in FHB of wheat and

barley. Wheat and *B. distachyon* have been show to share a common defence transcriptome

response following Fusarium crown rot infection caused by *F. pseudograminearum* (Powell

et al., 2017). Furthermore *B. distachyon* and barley were found to share common

brassinosteroid-associated defence responses through the use of brassinosteroid receptor

(*bri1*) mutants in the two species (Goddard *et al.*, 2014). This suggests the observed effects

of phytohormones on FHB in *B. distachyon* can be effectively translated to wheat FHB

responses.

The pathology of *F. graminearum* in root tissues is poorly characterised, but evidence

suggests that the fungus rapidly penetrates roots and exhibits specific stages of infection

(Wang *et al.*, 2015). Investigation of exogenous phytohormones on FRR response revealed

that FRR was much more responsive than FHB in the response to infection. This is likely due

to the continual exposure during FRR assays as opposed to the single application used for

FHB assays. An example of this was observed JA (Fig. 3B) and ACC (Fig. 3C) that displayed

the most pronounced effect on FHB susceptibility at earlier time points suggesting a short-

lived response.

The classic defence hormones SA, JA, and ethylene (ACC) exhibited different effects on

FRR and FHB (Fig. 5). Despite the non-significant FHB response to SA, SA increased FRR

susceptibility as early as 1 dpi (Fig. 1A). This difference may reflect the differences in the

levels of phytohormones in different tissues. Endogenous free SA levels were found to be

substantially higher in rice (*O. sativa*) floral tissues compared to root tissues (Chen *et al.*,

1997). If this is also true for *B. distachyon*, exogenous SA application might be expected to

have more impact on the endogenous SA profile in roots than florets resulting in a more

pronounced change in resistance in FRR compared to FHB. The similar response between JA

and ethylene in FHB and FRR supports the evidence that JA and ethylene often function

similarly in defence (Pieterse *et al.*, 2012). However, both JA and ethylene had contrasting

effects on FRR and FHB, increasing resistance to the former and susceptibility to the latter

(Fig. 1B-C and Fig 4B-C). The SA pathway is generally considered to promote resistance to

biotrophic pathogens and susceptibility to necrotrophic pathogens whereas JA and ethylene

pathways generally promote resistance to necrotrophic pathogens and susceptibility to

biotrophic pathogens (Glazebrook, 2005; Bari and Jones, 2009; Pieterse *et al.*, 2012). In both

FHB and FRR, hormones were applied before inoculation and would most likely influence

events relating to the biotrophic phase of *F. graminearum* infection and colonisation. Given

the increase susceptibility to FHB in response to JA and ethylene, this supports the JA/

ethylene pathway antagonism to the biotrophic phase of FHB. However, since the opposite

effect of JA and ethylene was observed on FRR, we speculate that *F. graminearum* is

adopting a more necrotrophic lifestyle on roots. This difference in trophism between tissues

is not well documented. Differences in trophic state have been observed for other plant

pathogens. Marcel and colleagues (2010) found that *M. oryzae* behaves as a biotroph

during rice root infection despite adopting a hemibiotrophic lifestyle on leaves (Marcel *et*

al., 2010). Hypothetical reasons for switching to necrotrophy may be due to a demand for a

change in the resource acquisition method, the time required to overcome defences, or a

change in plant defence response (Kabbage *et al.*, 2015; Zeilinger *et al.*, 2016). One or more

of these factors may be causing this difference in fungal lifestyle between roots and shoots

in the *B. distachyon - F. graminearum* pathosystem. Wang and colleagues (2015) found that

wheat Sumai3 roots were susceptible to FRR despite this variety being known for its potent

FHB resistance (Wang *et al.*, 2015). Furthermore, Lyons et al. (2015) identified differing

expression patterns between root and leaf infection by *Fusarium oxysporum*. These

examples suggest that a unique defence mechanism might be occurring in roots which

promotes a different trophic lifestyle in Fusarium.

420   Contrary to the classic defence hormones, the development-related phytohormones

induced similar effects on FHB and FRR (Fig. 5). Auxin application resulted in greatly

increased resistance in both FHB and FRR in *B. distachyon*. Like the effect of auxins, the

cytokinins trans-Zeatin and BABA also functioned in a tissue-independent manner, but they

greatly increased susceptibility to FHB and FRR (Fig. 5). Aside from roles in plant defence,

BABA can also affect plant development through reduction of growth and endogenous iron

content (Wu *et al.*, 2010; Koen *et al.*, 2014). Therefore, BABA could be considered to be

functioning similarly to the classic development-associated hormones auxin and cytokinin.

There was a substantially larger effect of these three phytohormones on resistance

compared to the classic defence phytohormones. We speculate that auxins, cytokinins, and

BABA are affecting resistance in an SA/JA independent manner. In support of this, (Qi *et al.*,

2016) found that exogenous application of IAA did not lead to an increase in endogenous

levels of SA or JA. Furthermore auxin signalling independent of JA and SA was found to be

important for *A. thaliana* resistance to two other necrotrophic pathogens (Llorente *et al.*,

2008). In another example, elevated expression of a the IAA-amido synthetase gene *GH3-8*

that regulates IAA levels promoted resistance to *X. oryzae* in rice independently of SA and JA

(Ding *et al.*, 2008). Cytokinins were also found to improve *N. tabacum* resistance to *P.*

syringae in an SA and JA independent manner (Großkinsky *et al.*, 2011). BABA functions

synergistically with the abscisic acid defence pathway (Ton and Mauch-Mani, 2004) and

there is evidence suggesting that abscisic acid promotes FHB susceptibility (Buhrow *et al.*,

2016; Qi *et al.*, 2016). The different effect of development hormones may be related to

either phytohormone homeostasis, antimicrobial production linked to metabolic pathways

of development hormones, or trade-offs with growth (Kazan and Manners, 2009; Huot *et*

al., 2014; Albrecht and Argueso, 2017). Additionally, the opposite and marked response

observed between cytokinins and auxins may be due to an antagonism between the two

response networks (Naseem and Dandekar, 2012; Naseem *et al.*, 2012). Overall, the effect

of development-associated hormones on FHB and FRR may depend on their influence on

basal resistance or specific growth/defence trade-offs independent of the trophic lifestyle.

In contrast, the role of the classic defence hormones may be dependent on the trophic

lifestyle of the pathogen.

Since multiple time points for both FHB and FRR were recorded, we observed differences

in slope regression lines between phytohormones. We were able to separate the responses

to compounds into two broad groups. The first 'parallel' group are those compounds that

induce a significant difference in resistance at the first time point, and that the difference

remained similar at later time points. The phytohormones SA, JA, and ACC (Fig. 1A-C) had

this effect on FRR whereas no phytohormone displayed this on FHB. This trend implies that

these compounds have an immediate effect on initial *F. graminearum* infection, after which

the absence of deviation relative to the first time point suggests that the effect on FHB

colonisation was much reduced or absent. The second 'divergent' group are those that have

minor effects at the first time point, but where the differential in resistance increased at

later time points. The growth and development-associated phytohormones exhibited this in

FRR assays: BABA (Fig. 1D), both auxins (Fig. 2A and Fig. 2B), and both cytokinins (Fig. 2C and

Fig. 2D). This trend implies that these compounds have a minimal effect on initial infection

but affect the ability of the host to restrict *F. graminearum* colonisation. These trends are

summarised in Fig. 5. Together the evidence supports the idea of fundamental differences in

the functioning of growth/development associated phytohormones compared to the classic

defence phytohormones in resistance to FRR. Secondly there were differences between FHB

and FRR in response to the classic defence hormone. This supports our view that for FHB

there is a period of biotrophic growth before switching to necrotrophy whereas in FRR this

initial phase is much reduced or absent.

A model was produced to summarise the effects of the tested phytohormones on FHB

and FRR in *B. distachyon* (Fig. 5). Phytohormones altered FRR and FHB resistance in *B.*

distachyon. The results reflect the generally antagonistic nature of SA to JA/ ethylene in the

two diseases. However they also show that these defence hormones induce disease-specific

effects between FHB and FRR. Unlike the classical defence-related hormones, the

development-associated hormones auxin and cytokinin had large tissue-independent effects

on *B. distachyon* response to FHB and FRR, highlighting an important role of these

hormones in the *B. distachyon* - *F. graminearum* pathosystem. Furthermore our data
suggest that each hormone functions at different phases with some affecting on initial *F.*
graminearum infection and others playing a greater role during colonisation. Future
experiments are required to examine the host and pathogen transcriptome response during
both FHB and FRR to provide additional evidence on the mode of trophism within infected
tissues and gene expression of both partners in the interaction.

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Conflicts of Interest

492   The authors declare no conflict of interest.

Supplementary Figures

<u>Fig. S1</u>: The growth (measured as radius from mycelial epicentre) of *F. graminearum* on
hormone-amended PDA at 2 dpa.

<u>Fig. S2</u>: The change in *F. graminearum*-induced FRR necrosis after application of gibberellic
acid (A) and epibrassinolide (B) on Bd3-1 seedling roots.

<u>Fig. S3</u>: The visual effect of 10 µM abscisic acid on 7-day old Bd3-1 roots in the absence of
disease.

<u>Fig. S4</u>: The visual decrease in RNL after IAA application (B) compared to control (A) at 6 dpi.

<u>Fig. S5</u>: The visual increase in RNL after Zeatin application (B) compared to DMSO control (A)
at 6 dpi.

<u>Fig. S6</u>: The change in number of *F. graminearum*-infected florets after four doses of salicylic
acid 24 h before, 30 min before inoculation, 3 dpi, and 7 dpi on Bd3-1.

<u>Fig. S.7</u>: *F. graminearum* PH1 mycelial growth on PDA after 3 dpa with 8 mM IAA and NAA
treated filter disks.

References

Abeles FB, Morgan PW, Saltveit Jr ME. 2012. *Ethylene in plant biology*: Academic press.

Albrecht T, Argueso CT. 2017. Should I fight or should I grow now? The role of cytokinins in plant growth and immunity and in the growth–defence trade-off. Annals of Botany **119**, 725-735.

Ali SS, Kumar GBS, Khan M, Doohan FM. 2013. Brassinosteroid Enhances Resistance to Fusarium Diseases of Barley. Phytopathology **103**, 1260-1267.

Ameye M, Audenaert K, De Zutter N, Steppe K, Van Meulebroek L, Vanhaecke L, De Vleesschauwer D, Haesaert G, Smagghe G. 2015. Priming of Wheat with the Green Leaf Volatile Z-3-Hexenyl Acetate Enhances Defense against Fusarium graminearum But Boosts Deoxynivalenol Production. Plant Physiology **167**, 1671-1684.

Bari R, Jones JDG. 2009. Role of plant hormones in plant defence responses. Plant Molecular Biology **69**, 473-488.

Beccari G, Covarelli L, Nicholson P. 2011. Infection processes and soft wheat response to root rot and crown rot caused by Fusarium culmorum. Plant Pathology **60**, 671-684.

Boddu J, Cho S, Kruger WM, Muehlbauer GJ. 2006. Transcriptome Analysis of the Barley-Fusarium graminearum Interaction. Molecular Plant-Microbe Interactions **19**, 407-417.

Brkljacic J, Grotewold E, Scholl R, Mockler T, Garvin DF, Vain P, Brutnell T, Sibout R, Bevan M, Budak H, Caicedo AL, Gao C, Gu Y, Hazen SP, Holt BF, Hong S-Y, Jordan M, Manzaneda AJ, Mitchell-Olds T, Mochida K, Mur LAJ, Park C-M, Sedbrook J, Watt M, Zheng SJ, Vogel JP. 2011. Brachypodium as a Model for the Grasses: Today and the Future. Plant Physiology **157**, 3-13.

Brown NA, Urban M, van de Meene AML, Hammond-Kosack KE. 2010. The infection biology of Fusarium graminearum: Defining the pathways of spikelet to spikelet colonisation in wheat ears. Fungal Biology **114**, 555-571.

Buhrow LM, Cram D, Tulpan D, Foroud NA, Loewen MC. 2016. Exogenous Abscisic Acid and Gibberellic Acid Elicit Opposing Effects on Fusarium graminearum Infection in Wheat. Phytopathology **106**, 986-996.

Chen X, Steed A, Travella S, Keller B, Nicholson P. 2009. Fusarium graminearum exploits ethylene signalling to colonize dicotyledonous and monocotyledonous plants. New Phytol **182**, 975-983.

Chen Z, Iyer S, Caplan A, Klessig DF, Fan B. 1997. Differential Accumulation of Salicylic Acid and Salicylic Acid-Sensitive Catalase in Different Rice Tissues. Plant Physiology **114**, 193-201.

Choi J, Choi D, Lee S, Ryu C-M, Hwang I. 2011. Cytokinins and plant immunity: old foes or new friends? Trends in Plant Science **16**, 388-394.

Choi J, Huh SU, Kojima M, Sakakibara H, Paek K-H, Hwang I. 2010. The Cytokinin-Activated Transcription Factor ARR2 Promotes Plant Immunity via TGA3/NPR1-Dependent Salicylic Acid Signaling in Arabidopsis. Developmental Cell **19**, 284-295.

Cohen Y. 2001. The BABA story of induced resistance. Phytoparasitica **29**, 375-378.

Cohen Y, Vaknin M, Mauch-Mani B. 2016. BABA-induced resistance: milestones along a 55-year journey. Phytoparasitica **44**, 513-538.

Cohen YR. 2002. β-Aminobutyric Acid-Induced Resistance Against Plant Pathogens. Plant Disease **86**, 448-457.

Cook RJ. 2001. Management of wheat and barley root diseases in modern farming systems. Australasian Plant Pathology **30**, 119-126.

Dean R, Van Kan JAL, Pretorius ZA, Hammond-Kosack KE, Di Pietro A, Spanu PD, Rudd JJ, Dickman M, Kahmann R, Ellis J, Foster GD. 2012. The Top 10 fungal pathogens in molecular plant pathology. Molecular Plant Pathology **13**, 414-430.

Ding X, Cao Y, Huang L, Zhao J, Xu C, Li X, Wang S. 2008. Activation of the Indole-3-Acetic Acid–Amido Synthetase GH3-8 Suppresses Expansin Expression and Promotes Salicylate- and Jasmonate-Independent Basal Immunity in Rice. The Plant Cell **20**, 228-240.

Draper J, Mur LAJ, Jenkins G, Ghosh-Biswas GC, Bablak P, Hasterok R, Routledge APM. 2001. Brachypodium distachyon. A New Model System for Functional Genomics in Grasses. Plant Physiology **127**, 1539-1555.

Foroud NA, Pordel R, Goyal RK, Ryabova D, Eranthodi A, Chatterton S, Kovalchuk I. 2018. Chemical activation of the ethylene signalling pathway promotes Fusarium graminearum resistance in detached wheat heads. Phytopathology.

Glazebrook J. 2005. Contrasting mechanisms of defense against biotrophic and necrotrophic pathogens. Annu. Rev. Phytopathol. **43**, 205-227.

Goddard R, Peraldi A, Ridout C, Nicholson P. 2014. Enhanced Disease Resistance Caused by BRI1 Mutation Is Conserved Between Brachypodium distachyon and Barley (Hordeum vulgare). Molecular Plant-Microbe Interactions **27**, 1095-1106.

Großkinsky D, Edelsbrunner K, Pfeifhofer H, van der Graaff E, Roitsch T. 2013. Cis- and trans-zeatin differentially modulate plant immunity. Plant Signaling & Behavior **8**, e24798.

Großkinsky DK, Naseem M, Abdelmohsen UR, Plickert N, Engelke T, Griebel T, Zeier J, Novák O, Strnad M, Pfeifhofer H. 2011. Cytokinins mediate resistance against Pseudomonas syringae in tobacco through increased antimicrobial phytoalexin synthesis independent of salicylic acid signaling. Plant Physiology **157**, 815-830.

Häffner E, Konietzki S, Diederichsen E. 2015. Keeping control: The role of senescence and development in plant pathogenesis and defense. Plants **4**, 449-488.

Huot B, Yao J, Montgomery BL, He SY. 2014. Growth–defense tradeoffs in plants: a balancing act to optimize fitness. Molecular Plant **7**, 1267-1287.

Jakab G, Cottier V, Toquin V, Rigoli G, Zimmerli L, Métraux J-P, Mauch-Mani B. 2001. β-Aminobutyric Acid-induced Resistance in Plants. European Journal of Plant Pathology **107**, 29-37.

Jansen C, von Wettstein D, Schäfer W, Kogel K-H, Felk A, Maier FJ. 2005. Infection patterns in barley and wheat spikes inoculated with wild-type and trichodiene synthase gene disrupted Fusarium graminearum. Proceedings of the National Academy of Sciences of the United States of America **102**, 16892-16897.

Kabbage M, Yarden O, Dickman MB. 2015. Pathogenic attributes of Sclerotinia sclerotiorum: switching from a biotrophic to necrotrophic lifestyle. Plant Science **233**, 53-60.

Kakei Y, Mochida K, Sakurai T, Yoshida T, Shinozaki K, Shimada Y. 2015. Transcriptome analysis of hormone-induced gene expression in Brachypodium distachyon. Scientific Reports **5**, 14476.

Kazan K, Manners JM. 2009. Linking development to defense: auxin in plant–pathogen interactions. Trends in Plant Science **14**, 373-382.

Kellogg EA. 2015. Brachypodium distachyon as a Genetic Model System. Annu Rev Genet **49**, 1-20.

Koen E, Trapet P, Brule D, Kulik A, Klinguer A, Atauri-Miranda L, Meunier-Prest R, Boni G, Glauser G, Mauch-Mani B, Wendehenne D, Besson-Bard A. 2014. beta-Aminobutyric Acid (BABA)-Induced Resistance in Arabidopsis thaliana: Link with Iron Homeostasis. Molecular Plant-Microbe Interactions **27**, 1226-1240.

Kouzai Y, Kimura M, Yamanaka Y, Watanabe M, Matsui H, Yamamoto M, Ichinose Y, Toyoda K, Onda Y, Mochida K, Noutoshi Y. 2016. Expression profiling of marker genes responsive to the defence-associated phytohormones salicylic acid, jasmonic acid and ethylene in Brachypodium distachyon. BMC Plant Biol **16**, 59.

Li G, Yen Y. 2008. Jasmonate and ethylene signaling pathway may mediate Fusarium head blight resistance in wheat. Crop Science **48**, 1888-1896.

Llorente F, Muskett P, Sánchez-Vallet A, López G, Ramos B, Sánchez-Rodríguez C, Jordá L, Parker J, Molina A. 2008. Repression of the Auxin Response Pathway Increases Arabidopsis Susceptibility to Necrotrophic Fungi. Molecular Plant **1**, 496-509.

Luo K, Rocheleau H, Qi PF, Zheng YL, Zhao HY, Ouellet T. 2016. Indole-3-acetic acid in Fusarium graminearum: Identification of biosynthetic pathways and characterization of physiological effects. Fungal Biology **120**, 1135-1145.

Makandar R, Essig JS, Schapaugh MA, Trick HN, Shah J. 2006. Genetically Engineered Resistance to Fusarium Head Blight in Wheat by Expression of Arabidopsis NPR1. Molecular Plant-Microbe Interactions **19**, 123-129.

Makandar R, Nalam V, Chaturvedi R, Jeannotte R, Sparks AA, Shah J. 2010. Involvement of Salicylate and Jasmonate Signaling Pathways in Arabidopsis Interaction with Fusarium graminearum. Molecular Plant-Microbe Interactions **23**, 861-870.

Makandar R, Nalam VJ, Lee H, Trick HN, Dong Y, Shah J. 2011. Salicylic Acid Regulates Basal Resistance to Fusarium Head Blight in Wheat. Molecular Plant-Microbe Interactions **25**, 431-439.

Mandal S, Mallick N, Mitra A. 2009. Salicylic acid-induced resistance to Fusarium oxysporum f. sp. lycopersici in tomato. Plant Physiology and Biochemistry **47**, 642-649.

Marcel S, Sawers R, Oakeley E, Angliker H, Paszkowski U. 2010. Tissue-adapted invasion strategies of the rice blast fungus Magnaporthe oryzae. The Plant Cell **22**, 3177-3187.

Mergoum M, Hill J, Quick J. 1998. Evaluation of resistance of winter wheat to Fusarium acuminatum by inoculation of seedling roots with single, germinated macroconidia. Plant Disease **82**, 300-302.

Miedaner T. 1997. Breeding wheat and rye for resistance to Fusarium diseases. Plant Breeding **116**, 201-220.

Naseem M, Dandekar T. 2012. The role of auxin-cytokinin antagonism in plant-pathogen interactions. PLoS pathogens **8**.

Naseem M, Philippi N, Hussain A, Wangorsch G, Ahmed N, Dandekar T. 2012. Integrated systems view on networking by hormones in Arabidopsis immunity reveals multiple crosstalk for cytokinin. The Plant Cell **24**, 1793-1814.

Opanowicz M, Hands P, Betts D, Parker ML, Toole GA, Mills ENC, Doonan JH, Drea S. 2011. Endosperm development in Brachypodium distachyon. Journal of Experimental Botany **62**, 735-748.

Pasquet JC, Chaouch S, Macadre C, Balzergue S, Huguet S, Martin-Magniette ML, Bellvert F, Deguercy X, Thareau V, Heintz D, Saindrenan P, Dufresne M. 2014. Differential gene expression and metabolomic analyses of Brachypodium distachyon infected by deoxynivalenol producing and non-producing strains of Fusarium graminearum. Bmc Genomics **15**, 17.

Peraldi A, Beccari G, Steed A, Nicholson P. 2011. Brachypodium distachyon: a new pathosystem to study Fusarium head blight and other Fusarium diseases of wheat. Bmc Plant Biology **11**, 100.

Petti C, Reiber K, Ali SS, Berney M, Doohan FM. 2012. Auxin as a player in the biocontrol of Fusarium head blight disease of barley and its potential as a disease control agent. Bmc Plant Biology **12**, 224-224.

Pieterse CM, Zamioudis C, Berendsen RL, Weller DM, Van Wees SC, Bakker PA. 2014. Induced systemic resistance by beneficial microbes. Annual Review of Phytopathology **52**, 347-375.

Pieterse CMJ, Does DVd, Zamioudis C, Leon-Reyes A, Wees SCMV. 2012. Hormonal Modulation of Plant Immunity. Annual Review of Cell and Developmental Biology **28**, 489-521.

Powell JJ, Carere J, Sablok G, Fitzgerald TL, Stiller J, Colgrave ML, Gardiner DM, Manners JM, Vogel JP, Henry RJ, Kazan K. 2017. Transcriptome analysis of Brachypodium during fungal pathogen infection reveals both shared and distinct defense responses with wheat. Scientific Reports **7**, 17212.

Qi P-F, Balcerzak M, Rocheleau H, Leung W, Wei Y-M, Zheng Y-L, Ouellet T. 2016. Jasmonic acid and abscisic acid play important roles in host–pathogen interaction between Fusarium graminearum and wheat during the early stages of fusarium head blight. Physiological and Molecular Plant Pathology **93**, 39-48.

Qi P-F, Johnston A, Balcerzak M, Rocheleau H, Harris LJ, Long X-Y, Wei Y-M, Zheng Y-L, Ouellet T. 2012. Effect of salicylic acid on Fusarium graminearum, the major causal agent of fusarium head blight in wheat. Fungal Biology **116**, 413-426.

Scholthof K-BG, Irigoyen S, Catalan P, Mandadi KK. 2018. Brachypodium: A Monocot Grass Model Genus for Plant Biology. The Plant Cell **30**, 1673-1694.

Sorahinobar M, Niknam V, Ebrahimzadeh H, Soltanloo H, Behmanesh M, Enferadi ST. 2016. Central Role of Salicylic Acid in Resistance of Wheat Against Fusarium graminearum. Journal of Plant Growth Regulation **35**, 477-491.

Sun Y, Xiao J, Jia X, Ke P, He L, Cao A, Wang H, Wu Y, Gao X, Wang X. 2016. The role of wheat jasmonic acid and ethylene pathways in response to Fusarium graminearum infection. Plant Growth Regulation **80**, 69-77.

Thevenet D, Pastor V, Baccelli I, Balmer A, Vallat A, Neier R, Glauser G, Mauch-Mani B. 2017. The priming molecule beta-aminobutyric acid is naturally present in plants and is induced by stress. New Phytologist **213**, 552-559.

Ton J, Mauch-Mani B. 2004. β-amino-butyric acid-induced resistance against necrotrophic pathogens is based on ABA-dependent priming for callose. The Plant Journal **38**, 119-130.

van Loon LC, Geraats BPJ, Linthorst HJM. 2006. Ethylene as a modulator of disease resistance in plants. Trends in Plant Science **11**, 184-191.

Vogel JP, Garvin DF, Leong OM, Hayden DM. 2006. Agrobacterium-mediated transformation and inbred line development in the model grass Brachypodium distachyon. Plant Cell, Tissue and Organ Culture **84**, 199-211.

Vogel JP, Garvin DF, Mockler TC, Schmutz J, Rokhsar D, Bevan MW, Barry K, Lucas S, Harmon-Smith M, Lail K. 2010. Genome sequencing and analysis of the model grass Brachypodium distachyon. Nature **463**, 763-768.

Wang Q, Vera Buxa S, Furch A, Friedt W, Gottwald S. 2015. Insights Into Triticum aestivum Seedling Root Rot Caused by Fusarium graminearum. Molecular Plant-Microbe Interactions **28**, 1288-1303.

Watt M, Schneebeli K, Dong P, Wilson IW. 2009. The shoot and root growth of Brachypodium and its potential as a model for wheat and other cereal crops. Functional Plant Biology **36**, 960-969.

Wu C-C, Singh P, Chen M-C, Zimmerli L. 2010. L-Glutamine inhibits beta-aminobutyric acid-induced stress resistance and priming in Arabidopsis. Journal of Experimental Botany **61**, 995-1002.

Zeilinger S, Gupta VK, Dahms TE, Silva RN, Singh HB, Upadhyay RS, Gomes EV, Tsui CK-M, Nayak S C. 2016. Friends or foes? Emerging insights from fungal interactions with plants. FEMS microbiology reviews **40**, 182-207.

Figure legends

Fig. 1: The change in *F. graminearum*-induced FRR necrosis after application of defence-related hormones 100 µM SA (**A**), 1 µM JA (**B**), 100 µM ACC (**C**), and 9 mM BABA (**D**) on Bd3-1 seedling roots. Each data point is the mean root necrosis length ± SE from two (**A**, **B**, **C**) or three (**D**) independent experiments, except score date 3 from **C** which was from one independent experiments. Score Date numbers (3-day intervals between each number) for **B** and **C** are the combined dpi from different experiments. Significance levels: ** p < .01, *** p < .001 compared to control.

Fig. 2: The change in *F. graminearum*-induced FRR necrosis after application of development-related auxins 10 µM IAA (**A**) and 5 µM IAA NAA (**B**) and cytokinins 10 µM trans-Zeatin (**C**) and 10 µM Kinetin (**D**) on Bd3-1 seedling roots. Each data point is the mean root necrosis length ± SE from three (**C**), two (**A** and **B**) or one (**D**) independent experiments. Significance level ** p < .01, *** p < .001 compared to control. Visual difference for **A** in Supp. Fig. 4 and **C** in Supp. Fig. 5.

Fig. 3: The change in number of *F. graminearum*-infected florets after pre-application of 200 µM SA (**A**), 50 µM JA (**B**), 1 mM ACC (**C**), 10 mM BABA, (**D**) and 10 µM trans-Zeatin (**E**) on Bd3-1. Each data point is the mean number of florets infected ± SE from one (**A**), two (**B**, **C**, **E**) or three (**D**) independent experiments. Score Date numbers (4-day intervals between each number) for **B** and **D** are the combined dpi from different experiments. Significance levels: * p < .05, ** p < .01, *** p < .001 compared to control.

Fig. 4: The number of *F. graminearum*-infected florets at 8 dpi after pre-application of IAA or NAA on Bd21. Each bar is the mean number of florets infected ± SE from one experimental replicate. Significance levels: ** p < .01, *** p < .001 compared to control.

Fig. 5: Model summarising all exogenous hormone application experiments (Fig. 1-5). Compares the effects between FHB and FRR in *B. distachyon* (on the left). This summary is a generalisation of the trend over time and includes different arrow thicknesses based on the potency of response caused by each phytohormone. Includes the predicted slope regression groups for each phytohormone presented as different arrow caps (excluding auxins and SA as these were not measured or not significantly different to control). Novel abbreviations: ET (ethylene), Aux (auxins), CK (cytokinins).

Figures
Fig. 1

Fig. 2

Fig. 3

A

B

C

D

E

Fig. 4

Fig. 5