



RESEARCH ARTICLE

Combination of yeast antagonists and Acibenzolar-S-Methyl reduced the severity of Fusarium head blight of wheat incited by *Fusarium graminearum sensu stricto*

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Abstract

The combination of yeast antagonists and Acibenzolar-S-Methyl (ASM) was tested against *Fusarium graminearum* on a spring wheat cultivar PAN3471. Two strains of *Papiliotrema flavescens* (Strains WL3 and WL6) and a strain of *Pseudozyma* sp. (MGO1) were combined with full strength ASM at anthesis, half strength ASM at anthesis and quarter strength ASM at late boot stages. The yeast and ASM treatments were applied prior to *F. graminearum* inoculation and disease progress was assessed over time. The combination of yeast and ASM treatments effectively reduced Fusarium Head Blight (FHB) severity and deoxynivalenol (DON) concentration compared to when the treatments were used alone. A positive correlation was observed between the Area Under Disease Progress Curve (AUDPC) and Percentage Seed Infection (PSI) ($r = 0.44$) whereas a negative correlation was observed between AUDPC and Hundred Seed Weight (HSW) ($r = -0.77$) and PSI and HSW ($r = -0.44$). The best combination treatment providing the highest reduction in final disease severity (41.83%), high HSW and moderate PSI was 0.075 g/l ASM at anthesis plus *P. flavescens* strain WL3. The highest DON reduction (19.35%) was by the treatment 0.075 g/l ASM at anthesis plus *P. flavescens* strain WL6. The best treatment was *P. flavescens* combined with 0.075 g/l ASM at anthesis. Although *Pseudozyma* sp. strain MGO1 did not provide the best FHB and DON reduction, its combination with ASM application improved disease control efficacy. To the best of our knowledge, this study presents the first report of the combination of *P. flavescens* and ASM in the management of FHB caused by *F. graminearum* in wheat plants.

Keywords

Biological control, induced resistance, *Papiliotrema flavescens*, *Pseudozyma* sp., mycotoxins, deoxynivalenol

Introduction

One of the major effects of Fusarium Head Blight (FHB) in crops is the production of mycotoxins in infected grains (1-3). These mycotoxins are a threat to human and animal health and have been reported to increase disease severity during infection by possibly disabling the plants' natural defence mechanisms (2, 4, 5). FHB infection is accompanied by the production of Fusarium-Damaged Kernels (FDKs) which cannot be used as either food, feed or seed (5, 6). The mycotoxins that are produced by *F. graminearum* [teleomorph *Gibberella zeae* (Schwein.) Petch], a predominant causal agent of FHB, are deoxynivalenol (DON) (and its derivatives), nivalenol (NIV) (and its derivatives) and zearalenone (ZEA) (5, 7, 8). Apart from DON being the

least harmful type B trichothecene mycotoxin, it is the most frequently detected mycotoxin and thus an indicator for mycotoxin contamination in FHB-infected grains (3, 4, 9). Previous studies indicate that pre-harvest control of FHB infection and development is the most promising means of reducing mycotoxin contamination on grains (5, 6, 10). This is because mycotoxin detoxification methods have limited efficacy on harvested grains and these methods have not yet been approved for use on grains with mycotoxin levels above acceptable limits (10).

Certain fungicides, such as triazole-based fungicides, have been used against FHB with reported efficacies (11-14). In resistance breeding programmes, some progress has been made which includes the identification of possible sources of resistance (such as the Chinese cultivar 'Sumai 3') (2, 5, 15). Although some FHB control efforts have shown potential in disease reduction, there are currently no registered fungicides or bio-fungicides, and no commercially available resistant wheat varieties in most parts of the world (2, 6, 16). Current research into the control of FHB has been aimed at the use of natural antagonists, resistance breeding and integrated management (2, 5, 6, 14, 17). These methods are of research interest since they address the issues associated with fungicide use which include chemical residues in/on food and the development of resistant pathogen strains (due to excessive use) (2, 4, 18). Biological control of plant diseases has been studied over the years with reported efficacy. Its advantages include reduced environmental hazards (compared to chemical use), reduced likelihood of resistance development, and the production of durable plant protection (19). BCAs can be applied on plant residues (13), plant tissue (13), soil (20, 21) and/or seed. Soil treated with *Trichoderma harzianum* and *T. viride* increased shoot dry weight, root dry weight and grain yield in the control of *Sclerotium rolfsii* (22). On the same study, the two *Trichoderma* species were reported to promote plant health by normalizing peroxidase (POX), phenylalanine ammonia lyase (PAL) and catalase (CAT) post inoculation with *S. rolfsii*. *Streptomyces* sp. RC 87B reduced FHB severity and DON by up to 39% and 85% respectively, on wheat during field trials (13). When applied on wheat stubble, *Streptomyces* sp. RC 87B reduced *F. graminearum* inoculum by at least 46% 90 days post inoculation (13).

It has been suggested that the best way to manage FHB is through integrated control strategies (11, 15, 23). Several studies on the incorporation of biological control agents (BCAs) in an integrated strategy for the control of FHB have been reported (24, 25). The co-culture of *Cryptococcus flavescens* OH 182.9 and *C. aureus* OH 71.4 significantly reduced FHB severity (by 32% on average) compared to individual applications (25). The integration of resistance inducers with BCAs in the management of FHB in wheat has been previously studied (26). However, this study reports for the first time the combination of Acibenzolar-S-Methyl (ASM) and yeast antagonists in an integrated management strategy of FHB caused by *F. graminearum*. The aim of this study was to test the efficacy of combining ASM with yeast antagonists for the reduction of FHB

severity and DON contamination in wheat. The effectiveness of the combined treatments as against each of the treatments alone was measured using the following parameters: (i) measure disease severity, (ii) Hundred Seed Weight (HSW), (iii) Percentage Seed Infection (PSI) and (iv) mycotoxin concentration (DON and ZEA) in harvested grains.

Materials and Methods

Planting and experimental design

Sixty-five planting pots of 25 cm diameter were filled up to 90% capacity with composted pine bark potting medium. Thereafter, spring wheat five seeds [cultivar PAN3471 obtained from Pannar Seed (Pvt) Ltd, Greytown, Republic of South Africa] were sown at even spacing in each pot which constituted an experimental unit. The trial consisted of 16 treatments (Table 1) with five replicates each. A completely randomised design was used for this experiment. The

Table 1. The description of the 16 treatments designed for the integration trial which consists of Acibenzolar-S-Methyl (ASM) and biological control agent (BCA) treatments.

Treatment no.	Treatment name	Description
1	ASM1	0.075 g/L ASM at anthesis (Feeke's 10.5.1)
2	ASM2	0.019 g/L ASM at late boot stage (Feeke's 10)
3	ASM3	0.0563 g/L ASM at anthesis (Feeke's 10.5.1)
4	BCA1	<i>Papiliotrema flavescens</i> WL3
5	BCA2	<i>Papiliotrema flavescens</i> WL6
6	BCA3	<i>Pseudozyma</i> sp. MGO1
7	ASM1 + BCA1	0.075 g/L ASM at anthesis (Feeke's 10.5.1) + <i>P. flavescens</i> WL3
8	ASM1 + BCA2	0.075 g/L ASM at anthesis (Feeke's 10.5.1) + <i>P. flavescens</i> WL6
9	ASM1 + BCA3	0.075 g/L ASM at anthesis (Feeke's 10.5.1) + <i>Pseudozyma</i> sp. MGO1
10	ASM2 + BCA1	0.019 g/L ASM at late boot stage (Feeke's 10) + <i>P. flavescens</i> WL3
11	ASM2 + BCA2	0.019 g/L ASM at late boot stage (Feeke's 10) + <i>P. flavescens</i> WL6
12	ASM2 + BCA3	0.019 g/L ASM at late boot stage (Feeke's 10) + <i>Pseudozyma</i> sp. MGO1
13	ASM3 + BCA1	0.0563 g/L ASM at anthesis (Feeke's 10.5.1) + <i>P. flavescens</i> WL3
14	ASM3 + BCA2	0.0563 g/L ASM at anthesis (Feeke's 10.5.1) + <i>P. flavescens</i> WL6
15	ASM3 + BCA3	0.0563 g/L ASM at anthesis (Feeke's 10.5.1) + <i>Pseudozyma</i> sp. MGO1
16	Control	No ASM and no yeast antagonist

Key: Feeke's 10 and Feeke's 10.5.1 are wheat growth stages presented by the Feeke's scale (28).

pots were placed in a growing area with insect netting (approximately 15% shading) and a drip irrigation system was used where each pot received water for 2 mins four times a day. Osmocote Exact Mini 5-6 M 15-3.9-9.1 + 1.2 Mg + TE [supplied by Greenhouse products (Pvt) Ltd, Helderkruin, Republic of South Africa (RSA)], an ammonium based slow-release fertilizer, was applied in each pot at a rate of 2.5 g/l of potting media.

Inoculum preparation

A *F. graminearum* (strain F20) conidia suspension previously stored at -80 °C, was thawed under a laminar flow cabinet at ambient temperature. This strain was obtained from the Discipline of Plant Pathology stock culture laboratory, University of KwaZulu-Natal, Pietermaritzburg, South Africa. The *F. graminearum* strain was previously isolated from infected wheat heads in a wheat cultivation field. Conidia were then streaked out onto fresh potato dextrose agar (PDA) plates and incubated at 25 °C for 5 days. Thereafter, the culture was subcultured by cutting out a 1 mm³ agar plug from the actively growing edges of the mycelia and then placed faced down at the centre of a fresh PDA plate. This was repeated on 20 PDA plates and the plates were incubated at 25 °C for 7 days. Thereafter, the plates were placed under ultraviolet-A (UVA) light (360 nm wavelength) for 14 days to induce fungal sporulation.

The yeast strains *Papiliotrema flavescens* [strains WL3 and WL6, previously isolated from wheat (*Triticum aestivum* L.) leaves] and *Pseudozyma* sp. [strain MGO1 previously isolated from Mondo grass (*Ophiopogon japonicus* (L.f.) Ker-Gaw) leaves] with proven efficacy against *F. graminearum* *in vitro* (27) were used in this study. These strains had been identified by Inqaba Biotechnological industries (Pvt) Ltd (Muckleneuk, Pretoria, RSA) using Internal Transcribed Spacers (ITS) sequencing and molecular identification. The yeast strains were streaked out from their respective stock solutions (previously stored at -80 °C) onto fresh PDA plates with 10 replicates each and thereafter incubated at 25 °C for 5 days. Thereafter, 4 ml of sterile distilled water was pipetted onto each plate using a micropipette under aseptic conditions. Using a flame-sterilized L-bent glass rod, the culture was suspended in the water by lightly rubbing the surface of the plate. The aliquot was decanted into a sterile and appropriately labelled conical flask. This was repeated for all the plates resulting in three flasks containing each yeast isolate.

Conidial suspensions of *F. graminearum* F 20 were prepared as above. The aliquot was transferred into a sterile Schott bottle which was vigorously shaken to allow the suspension of conidia in the solution. The aliquot was sieved through a sterile cheesecloth to remove mycelia and agar debris. Thereafter, the conidial concentration was adjusted to 1×10⁵ conidia/ml using a haemocytometer and then made up to 10 l. The spore concentrations of each of the 3 yeasts were adjusted to 1×10⁷ spores/ml and the solutions were made up to 4 l each.

Treatments application

Acibenzolar-S-Methyl (ASM) granules were purchased from Syngenta (Pvt) Ltd, Halfway house, Johannesburg, RSA. To prepare ASM concentrations, beakers were filled with tap water and placed on a bench top for an hour to allow the release of excess chlorine. ASM granules were weighed (0.019 g, 0.0563 g and 0.075 g) and each amount separately dissolved in 1 L of the tap water. The ASM solutions were transferred to previously cleaned and appropriately labelled 1 l pump spray bottles. With the nozzle adjusted to emit a fine mist, the plants were sprayed with the appropri-

ate ASM solutions until runoff at the appropriate growth stages (Table 1).

Wheat heads were sprayed with the appropriate yeast spore suspensions until runoff according to the assigned treatments presented in Table 1. In all *in vivo* inoculations, plants of the same treatment were sprayed separately, away from the other plants to prevent spray drift. The heads were then covered with perforated, light-weight plastic bags for 24 hrs. to encourage humidity. Forty-eight hrs after yeast inoculation, the wheat heads were sprayed with conidial suspensions of *F. graminearum* until runoff and thereafter covered with the same plastic bags for 24 hrs to encourage disease development. Yeast and *F. graminearum* inoculations were each performed once. Disease severity was measured using a visual scale originally described by (29) and disease ratings were recorded in intervals over time. The experiment was repeated once.

When the plants had a golden-brown appearance and had reached maturity, wheat heads were cut off from the straws and put in appropriately labelled collection bags according to treatment replicates. Harvested grains were placed in a ventilating oven set at 55°C for a period of 4 days. During this period, the bags were constantly monitored and shuffled to prevent heat damage of the grains. Thereafter, the wheat heads were threshed, and the seeds were transferred into appropriately labelled envelopes. These were stored in a cold room set at 4°C for further experiments.

HSW and PSI were determined per treatment replicate for the 2 experiments. For the PSI, the seeds were surface sterilized, cultured on freshly prepared PDA plates and incubated at 25°C for 4 days. Since each treatment had 5 replicates, each replicate had 3 plates which each had 15 seeds. The experiment was repeated once resulting in 30 plates per treatment. The number of Fusarium-infected seeds per plate was recorded and used to calculate the PSI using the following formula:

$$\text{Percentage Seed Infection (PSI)} = \frac{\text{No. of infected seeds}}{\text{Total seeds plated}} \times 100$$

Mycotoxin analysis

The target mycotoxins were DON and ZEA since they are the most prevalent mycotoxins in FHB infections. The roQ™ QuEChERS kits KSO-8909 and KSO-9507 were used for sample extraction and dispersive Solid Phase Extraction (dSPE) respectively. These were purchased from Separations (Pvt) Ltd, Johannesburg, RSA. Mycotoxin extraction was performed according to (30), with modifications. Wheat seeds from the 2 experiments were pooled according to treatments for mycotoxin analysis. For each sample, the seeds were ground into fine powder using a Mikro-Feinmuhle-Cullati (MFC) plant grinder in the Plant Pathology seeds laboratory. A 5 g subsample was added into a 50 ml roQ QuEChERS extraction tube along with the following reagents: Milli-Q water (10 ml), acetonitrile with 5% formic acid (10 ml) and the contents of the roQ QuEChERS extraction packet (KSO-8909) which consisted of 4.0g MgSO₄, 1.0 g NaCl, 1.0 g SCTD and 0.5 g SCDS (30).

The tube was shaken for 1 min by hand and then centrifuged at 4000 rpm (3000 g) for 5 mins (Beckman Coulter®, Avanti® J-26 XPI centrifuge) (30). Six ml (6 ml) of the supernatant were transferred into a roQ QuEChERS 15 ml centrifuge tube (KSO-9507) containing 900 mg MgSO₄ and 150 mg primary secondary amine (PSA) (30). The tube was shaken by hand for 30 secs and then centrifuged as above (30). Thereafter, 1 ml of the supernatant was filtered through a 0.45 µm pore filter and transferred into a 1.5 ml autosampler vial (all purchased from Separations (Pvt) Ltd, Johannesburg, Republic of South Africa) (30). This was repeated for all the samples and the vials were left open in a laminar flow cabinet overnight to dry.

The samples were analysed for the quantification of DON and ZEA using High Performance Liquid Chromatography (HPLC) (30). The HPLC system consisted of an LC-2030 pump connected to an LC-2030/2040 PDA detector, LC-2030 controller and LC-2030 autosampler. Chromatographic separations were performed on a Kinetex® 5 µm Biphenyl 100A LC Column (100 x 2.1 mm) connected to a guard column SecurityGuard™ filled with the same phase (2.1 to 4.6 mm) (all purchased from Separations (Pvt) Ltd, Johannesburg, Republic of South Africa). The mobile phase consisted of aqueous 5 mM ammonium acetate with 0.1% acetic acid, 5 mM ammonium acetate in methanol with 0.1% acetic acid, acetonitrile and Milli-Q water. The flow rate was 0.20 ml/min and the injection volume was 50 µl. The retention times for DON and ZEA were 2.563 and 10.193 mins respectively. Quantification was relative to external standards of 1–8 µg/ml in acetonitrile. Three quantification readings were conducted per sample.

Data analysis

HSW, PSI and disease severity data obtained were checked for homogeneity within the repeated trials and the data were thereafter pooled. Disease severity data was used to calculate the Area Under the Disease Progress Curve (AUDPC) for all treatments (31) before subjected to ANOVA. If the ANOVA was significant ($P \leq 0.05$), the means were separated using the Duncan's Multiple Range Test (DMRT) at 5% significance level using SAS software Version 9.4 (32). Pairwise correlations were determined between AUDPC, HSW and PSI for the pooled data using the Spearman's correlation test (32). The rate of disease progress (r) was calculated using the Vanderplank's logistic equation (33) expressed below:

$$r = \frac{1}{t_2 - t_1} \left[\left(\ln \frac{x_2}{1 - x_2} \right) - \left(\ln \frac{x_1}{1 - x_1} \right) \right]$$

where; t_1 = initial day of rating; t_2 = final day of rating; x_1 = initial disease value; x_2 = final disease value.

Results

Disease severity and seed infection studies

The 0.075 g/l ASM treatment at anthesis plus *P. flavescens* WL6 had the lowest disease severity rating in all rating days and thus the lowest final average disease severity (50.92%) compared to the control (87.53%) (Table 2). This means that the number of infected spikes for the treatment 0.075 g/l ASM at anthesis plus *P. flavescens* WL6 were significantly less than those for the control treatment. The control treatment had the highest average disease severity

Table 2. The average Fusarium Head Blight (FHB) severities of the 16 treatments throughout the disease rating days

Treatment No.	Treatment name	Number of days (Disease Rating days)							
		0	15	19	22	25	29	33	43
1	0.075 g/L ASM at anthesis	0	22.83	31.75	36.75	41.33	46.13	51.75	52.38
2	0.019 g/L ASM at late boot stage	0	27.58	38.62	44.46	56.33	63.63	68.31	73.94
3	0.0563 g/L ASM at anthesis	0	31.83	39.25	44.88	54.25	56.75	60.08	64.88
4	<i>P. flavescens</i> WL3	0	34.04	44.88	51.54	58.42	68.63	70.71	73.00
5	<i>P. flavescens</i> WL6	0	20.75	34.04	38.88	44.50	49.46	54.15	56.54
6	<i>Pseudozyma</i> sp. MGO1	0	15.29	37.38	42.38	49.87	55.71	62.58	69.35
7	0.075 g/L ASM at anthesis plus <i>P. flavescens</i> WL3	0	21.21	31.96	39.25	47.17	54.88	59.36	63.10
8	0.075 g/L ASM at anthesis plus <i>P. flavescens</i> WL6	0	10.80	24.08	27.71	36.33	41.96	50.08	50.92
9	0.075 g/L ASM at anthesis plus <i>Pseudozyma</i> sp. MGO1	0	16.29	30.42	36.96	44.15	48.83	57.79	59.88
10	0.019 g/L ASM at late boot stage plus <i>P. flavescens</i> WL3	0	22.50	32.38	39.67	47.58	51.54	56.96	62.38
11	0.019 g/L ASM at late boot stage plus <i>P. flavescens</i> WL6	0	22.38	32.06	35.71	47.79	51.75	58.21	59.46
12	0.019 g/L ASM at late boot stage plus <i>Pseudozyma</i> sp. MGO1	0	19.17	34.88	38.00	49.46	53.42	59.25	67.27
13	0.0563 g/L ASM at anthesis plus <i>P. flavescens</i> WL3	0	14.63	27.04	33.42	42.58	48.00	50.50	59.46
14	0.0563 g/L ASM at anthesis plus <i>P. flavescens</i> WL6	0	15.38	28.63	38.83	47.38	52.63	55.71	60.92
15	0.0563 g/L ASM at anthesis plus <i>Pseudozyma</i> sp. MGO1	0	22.48	36.54	43.21	55.92	61.13	63.63	69.88
16	Control	0	49.92	60.48	66.69	74.62	79.88	83.00	87.53
Effects		F-value							P-value
Treatments		30.5							0.0001
Time (Days)		492.43							0.0001
Treatments*Time (Days)		0.95							0.633
%CV									26.7

in all rating days (Table 2).

There were significant differences between the treatments for the AUDPC, HSW and PSI at $p \leq 0.0006$, $p \leq 0.0001$ and $p \leq 0.06$ respectively (Table 3). The lowest AUDPC and PSI values were observed for treatments 0.075 g/l ASM at anthesis plus *P. flavescens* WL6 and 0.075 g/l ASM at anthesis plus *Pseudozyma* sp. MGO1 respectively. The highest HSW was observed for the treatment 0.0563 g/l ASM at anthesis plus *P. flavescens* WL3. The PSI values of 8 out of 15 treatments were not significantly different from

nificant at $p = 0.0002$. A strong and moderate negative correlation was observed between AUDPC and HSW ($r = -0.77$) and PSI and HSW ($r = -0.44$) respectively. These were significant at $p < 0.0001$ and $p = 0.0003$, respectively. The highest negative correlation was observed between AUDPC and HSW.

Mycotoxin analysis

A reduction of up to 19.45% in DON concentration was obtained and this was by the treatment 0.075 g/l ASM at anthesis plus *P. flavescens* WL3 (Table 5). Although statistical-

Table 3. The average Area Under the Disease Progress Curve (AUDPC), average Hundred Seed Weight (HSW) (g), average Percentage Seed Infection (PSI) (%) and rate of disease progress (r) for the 16 treatments tested against Fusarium Head Blight (FHB) *in vivo*

Treatment	AUDPC	Average HSW (g)	Average PSI (%)	(r)
0.075 g/L ASM at anthesis	1113.27±122.90 ^{cd}	2.96±0.23 ^{ab}	39.58±12.02 ^{bc}	0.06
0.019 g/L ASM at late boot stage	1830.14±127.38 ^{bc}	2.26±0.15 ^b	61.11±12.58 ^{ab}	0.08
0.0563 g/L ASM at anthesis	1389.00±182.12 ^{bc}	2.81±0.24 ^{ab}	54.17±5.13 ^{abc}	0.05
<i>Papiliotrema flavescens</i> WL3	1579.20±60.66 ^b	1.86±0.21 ^c	52.78±4.39 ^{abc}	0.06
<i>Papiliotrema flavescens</i> WL6	1158.57±118.53 ^{bcd}	2.92±0.23 ^{ab}	34.72±1.79 ^{bc}	0.07
<i>Pseudozyma</i> sp. MGO1	1268.36±111.00 ^{bcd}	2.79±0.23 ^{ab}	56.94±8.67 ^{abc}	0.10
0.075 g/L ASM at anthesis plus <i>P. flavescens</i> WL3	1237.34±167.87 ^{bcd}	2.85±0.29 ^{ab}	37.50±8.06 ^{bc}	0.08
0.075 g/L ASM at anthesis plus <i>P. flavescens</i> WL6	936.16±182.93 ^d	3.32±0.22 ^{ab}	49.31±2.63 ^{abc}	0.09
0.075 g/L ASM at anthesis plus <i>Pseudozyma</i> sp. MGO1	1140.69±154.95 ^{bcd}	3.18±0.19 ^{ab}	31.94±8.60 ^c	0.08
0.019 g/L ASM at late boot stage plus <i>P. flavescens</i> WL3	1223.48±88.46 ^{bcd}	3.06±0.10 ^{ab}	51.39±10.42 ^{abc}	0.07
0.019 g/L ASM at late boot stage plus <i>P. flavescens</i> WL6	1208.75±146.97 ^{bcd}	3.11±0.22 ^{ab}	41.67±9.00 ^{bc}	0.06
0.019 g/L ASM at late boot stage plus <i>Pseudozyma</i> sp. MGO1	1244.82±134.54 ^{bcd}	3.01±0.22 ^{ab}	50.69±2.86 ^{abc}	0.09
0.0563 g/L ASM at anthesis plus <i>P. flavescens</i> WL3	1060.53±156.16 ^{cd}	3.41±0.26 ^a	36.11±11.05 ^{bc}	0.09
0.0563 g/L ASM at anthesis plus <i>P. flavescens</i> WL6	1146.89±85.97 ^{bcd}	3.34±0.18 ^{ab}	50.69±9.17 ^{abc}	0.08
0.0563 g/L ASM at anthesis plus <i>Pseudozyma</i> sp. MGO1	1364.83±124.09 ^{bcd}	3.01±0.13 ^{ab}	44.44±4.09 ^{bc}	0.08
Control	1964.98±97.08 ^a	1.77±0.23 ^c	72.92±6.84 ^a	0.07
F value	3.42	4.30	1.82	
P value	0.0006	<0.0001	0.06	
%C.V.	20.67	22.89	33.69	

Values followed by the same superscript letter are statistically identical

the control (Table 3). As a result, some treatments with low AUDPC and HSW values were associated with high PSI values. An example of this was 0.075 g/l ASM at anthesis plus *P. flavescens* WL6, which had the lowest AUDPC units, the third highest HSW and a below-average PSI that was not significantly different from the control. The control had the highest AUDPC units, lowest HSW and highest PSI. The highest rate of disease progress (r) was observed for *Pseudozyma* sp. MGO1 and the lowest for 0.0563 g/l ASM at anthesis (Table 3). However, there was not much difference in the rate of disease progress between treatments and thus statistical analysis was omitted.

Correlation between AUDPC, HSW and PSI

Significant correlations were observed for all pairwise combinations (Table 4). A moderate positive correlation was observed between AUDPC and PSI ($r = 0.44$) which was sig-

Table 4. Pairwise correlation between the Area Under the Disease Progress Curve (AUDPC), Hundred Seed Weight (HSW) and Percentage Seed Infection (PSI) for the 16 treatments tested against Fusarium Head Blight.

	AUDPC	HSW	PSI
AUDPC	1	-0.77*	0.44**
HSW		1	-0.44***
PSI			1

Key: (*) = $p < 0.0001$; (**) = $p = 0.0002$; (***) = $p = 0.0003$

ly similar to the DON concentration of the control, the other treatments with low DON concentrations were 0.075 g/l ASM at anthesis and 0.075 g/l ASM at anthesis plus *P. flavescens* WL6 respectively. The treatment 0.019 g/l ASM at late boot stage was the only treatment that had a DON concentration higher than that of the control. ZEA was not detected in all tested treatments and the control, as indicated

Table 5. Concentrations of deoxynivalenol (DON) and zearalenone (ZEA) in harvested grains for the 15 treatments compared to the control.

Treatment	DON		ZEA	
	Concentration (ppm)	Percentage reduction (%)	Concentration (ppm)	Percentage reduction (%)
0.075 g/L ASM at anthesis	12.45 ^{ab}	14.69	0.00	0.00
0.019 g/L ASM at late boot stage	15.39 ^a	*	0.00	0.00
0.0563 g/L ASM at anthesis	13.74 ^{ab}	5.82	0.00	0.00
<i>P. flavescens</i> WL3	13.28 ^{ab}	9.01	0.00	0.00
<i>P. flavescens</i> WL6	13.94 ^{ab}	4.48	0.00	0.00
<i>Pseudozyma</i> sp. MGO1	14.07 ^{ab}	3.54	0.00	0.00
0.075 g/L ASM at anthesis plus <i>P. flavescens</i> WL3	11.77 ^b	19.35	0.00	0.00
0.075 g/L ASM at anthesis plus <i>P. flavescens</i> WL6	12.76 ^{ab}	12.54	0.00	0.00
0.075 g/L ASM at anthesis plus <i>Pseudozyma</i> sp. MGO1	12.94 ^{ab}	11.28	0.00	0.00
0.019 g/L ASM at late boot stage plus <i>P. flavescens</i> WL3	13.25 ^{ab}	9.20	0.00	0.00
0.019 g/L ASM at late boot stage plus <i>P. flavescens</i> WL6	14.21 ^{ab}	2.61	0.00	0.00
0.019 g/L ASM at late boot stage plus <i>Pseudozyma</i> sp. MGO1	13.25 ^{ab}	9.18	0.00	0.00
0.0563 g/L ASM at anthesis plus <i>P. flavescens</i> WL3	13.12 ^{ab}	10.05	0.00	0.00
0.0563 g/L ASM at anthesis plus <i>P. flavescens</i> WL6	13.53 ^{ab}	7.29	0.00	0.00
0.0563 g/L ASM at anthesis plus <i>Pseudozyma</i> sp. MGO1	13.02 ^{ab}	10.76	0.00	0.00
Control	14.59 ^{ab}	0.00	0.00	0.00
F value	1.53		0.00	
P value	0.13		0.00	
%C.V.	13.09		0.00	

Key: (*) = values less than zero (negative values) indicating an increase in DON concentration compared to the control. Values followed by the same superscript letter are statistically identical

by 0.00 ppm in Table 5.

Discussion

In this study, we demonstrated that the combination of a plant defence inducer (ASM) with yeast antagonist reduced the severity of FHB and DON concentration in wheat. There is limited research on the integration of plant defence inducers with yeast antagonists in the management of *F. graminearum* in wheat. However, our results reveal good potential for FHB and DON reduction in wheat. When ASM was applied alone, the treatment 0.075 g/l ASM at anthesis had the lowest AUDPC units (1113.27), highest HSW (2.96 g) and the lowest DON concentration (12.45 ppm). When the biocontrol agents were applied alone, the treatment *P. flavescens* WL6 had the lowest AUDPC units (1158.57) and highest HSW (2.92 g), but *P. flavescens* WL3 had the lowest DON concentration (13.28 ppm). These treatments were previously tested prior to the current experiment (27) where their potential was studied. Although there is not much literature on the study of ASM for FHB control, ASM has been tested on other plant pathogens (34, 35). In a study to test the effect of ASM against *Botrytis cinera* on table grapes, ASM reduced the incidence of gray mold by up to 85% (34). Yeast biocontrol agents such as *Cryptococcus flavescens* OH 182.9 have been studied and reported to effectively reduce FHB and DON concentrations in wheat (24–26).

The integration of ASM and biocontrol agents improved FHB reduction compared to when the treatments were applied alone. The best integration treatments providing the highest FHB and DON reduction were 0.075 g/

l ASM at anthesis plus *P. flavescens* WL6 (41.83%) and 0.075 g/l ASM at anthesis plus *P. flavescens* WL3 (19.35%) respectively. Moreover, the treatment 0.075 g/l ASM at anthesis plus *P. flavescens* WL6 had the highest reduction in AUDPC units (52.91%), high DON reduction (12.54%), high HSW (3.32 g) and a PSI below 50%. It is important to note that the best integration treatments were comprised of treatments that performed best amongst those applied alone, which was expected. In another study, the combination of 4 resistance inducers with the yeast antagonist *Cryptococcus flavescens* OH 182.9 did not significantly reduce FHB severity compared to when applied alone (24). However, lowest FHB severity values were often associated with integrated treatments (24). Our study, therefore, is the first to report effective reduction of FHB and DON in wheat following the integration of ASM treatments and yeast biocontrol agents.

An increase in FHB severity is accompanied by an increase in *Fusarium*-infected kernels and a reduction in seed weight (4, 36). Similar correlations were also observed in our study thus aiding to the efficacy of these treatments in FHB management. Although majority the PSI values in our study were not significantly different to that of the control, the PSI of the tested treatments could potentially decrease with higher treatment application doses or integration with postharvest control methods. *Pseudozyma* sp. MGO1, which was isolated from the weed plant *Ophiopogon japonicus* (Mondo grass), although did not provide the best FHB and DON reduction compared to *P. flavescens* WL3 and *P. flavescens* WL6 treatments, the combination with ASM application resulted in increased efficacy. The treatment 0.075 g/l ASM at anthesis plus *Pseudozyma* sp. MGO1 was the best treatment amongst those treated with *Pseudozy-*

ma sp. MGO1, providing a high DON reduction of 11.28% compared to other treatments. *O. japonicus* is a traditional Chinese medicinal plant with antifungal activities. There is currently no literature available on its antifungal properties being tested against *F. graminearum*. Our previous study (27) contains *in vitro* and *in vivo* screening experiments on the antifungal effect of yeast isolates (such as MGO1) isolated from *O. japonicus* against *F. graminearum*. Therefore, this is novel work which shows potential and requires more research.

Although the two *P. flavescens* strains belong to the same species, the differences in their efficacies against *F. graminearum* shows that they could be different strains. Nevertheless, the combinations of the *P. flavescens* WL6 with 0.075 g/l ASM at anthesis was the overall best treatment in this study. Moreover, further research on the determination of the best inoculum dosage, frequency of application is required and could help improve the efficacy and reliability of the *P. flavescens* strains. Other studies in the control of plant diseases include testing ASM concentrations higher than the ones used in this study (34, 35). Therefore, future research can be aimed at determining the efficacy of higher ASM concentrations in the control of FHB of wheat as well as the physiological effects of the treatments on the wheat plant.

Conclusion

This study provides extensive research into the efficacy of Acibenzolar-S-Methyl (ASM) in the integrated control of Fusarium Head Blight (FHB) incited by *F. graminearum* in wheat plants. The combination of ASM and the *P. flavescens* strains provided the best FHB and deoxynivalenol (DON) reduction compared to when either were applied alone. The highest reduction in final FHB severity and DON concentration was observed where 0.075g/l ASM was applied at anthesis in combination with *P. flavescens* strains WL6 and WL3 respectively. To the best of our knowledge, this study presents the first report of *P. flavescens* strains as combination treatments with ASM in the management of FHB caused by *F. graminearum* in wheat plants. Field studies are essential to determine the efficacy of combined use of ASM and the *P. flavescens* strains in environments similar to those present in commercial wheat cultivation systems.

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Authors contributions

SS carried out the work and wrote the initial draft of the manuscript, implemented the comments after editing and revised the manuscript. NM – student co-supervision, funding acquisition and proof-reading of the final draft. KSY – student supervision, project leadership and management, funding acquisition and editing of manuscript.

Compliance with ethical standards

Conflict of interest: Authors do not have any conflict of interests to declare.

Ethical issues: None.

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