28 Minutes Later: A behavioural and transcriptomic investigation into the potential role of

aflatrem-like compounds in establishing *Ophiocordyceps* extended phenotypes in zombie

ants

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ABSTRACT

Coevolutionary relationships between parasites and their hosts can lead to bizarre phenotypes over

time. As such, certain *Ophiocordyceps* fungi have evolved to manipulate the behaviour of insect

and arachnid hosts. The most conspicuous examples are found in ants of the Camponotini tribe,

colloquially known as "zombie ants", which climb to elevated positions to aid fungal spore

transmission. While the unusual behaviours induced during infection are well described, the

molecular driving force behind these changes is still unknown. Recent genomics and

transcriptomics analyses have identified several candidate secreted proteins and secondary

metabolites, highly upregulated during infection. Amongst these compounds is an ergot alkaloid

related to the mycotoxin aflatrem, known to cause neurological issues in cows collectively called

"staggers syndrome". Staggering is also observed in *Camponotus floridanus* ants during the late

stages of *Ophiocordyceps* infection and manipulation. To test if aflatrem-like compounds could be responsible for this behavioural phenotype, we injected healthy ants with purified aflatrem, and recorded their behaviour for 30 minutes. Using both the machine learning tool MARGO and manual behavioural quantification, we found that aflatrem caused reduced ant activity and speed, and increased the number and duration of staggering behaviours. To examine changes at the transcriptome level that might underlie these phenotypes, we performed RNA-Seq on the heads of aflatrem-injected ants. We identified 261 genes that were significantly up- or down-regulated in the aflatrem-injected ants compared to sham-injected controls. When compared with RNA-seq data available for ants manipulated by *Ophiocordyceps camponoti-floridani*, we found that 108 of these genes were also differentially regulated in the heads of late-stage *Ophiocordyceps*-infected ants. These shared dysregulated genes included *sensory neuron membrane protein* genes, several *odorant-binding protein* genes, and musculoskeletal genes such as *titin* and *obscurin*. Together, these results indicate that aflatrem-like compounds, produced by *O. camponoti-floridani* during behavioural manipulation, significantly affect neuromuscular and sensory functioning, leading to a staggers phenotype in *C. floridanus*.

1. INTRODUCTION

As parasites adapt to their hosts, the two often develop an intense relationship driven by selective pressures presented in the host environment. As the parasite becomes more successful, the selective burdens shift to favour emergent resistance phenotypes in the host. Over time, these co-evolutionary pressures can lead to host-specificity and the subsequent emergence of unique manipulation phenotypes. Host manipulations provide an adaptive advantage to the parasite, for instance through the induction of specific behaviours that increase its reproduction success

(Sánches and Biron, 2019). These adaptive phenotypes are different from sickness behaviours,

defined as actions exhibited by infected individuals that are not conducive to the fitness of the host

nor the parasite (e.g., lethargy). One example of behavioural manipulation can be observed in the

parasitic flat worm *Leucochloridium paradoxum*, which manipulates amber snails from the genus

Succinea. Once inside a snail, the worm drives the host to ascend nearby structures, a common

manipulation behaviour known as "summit disease" (Wesołowska and Wesołowski, 2013).

Summiting behaviour makes snails more susceptible to predatory birds, the primary host of *L.*

paradoxum, ultimately increasing the parasite's likelihood of completing its lifecycle

(Wesołowska and Wesołowski, 2013).

Such behaviour-manipulating parasites are often colloquially referred to as "zombie

parasites" because the phenotypes exhibited by their hosts are reminiscent of zombies from pop

culture and sci-fi media (e.g., the movies *28 Days Later* and *The Body Snatchers*). While the

manifestations of behavioural manipulation vary across species, many host-manipulating parasites

have convergently evolved to induce summiting behaviour to increase their transmission (Araújo

& Hughes, 2019; de Bekker et al., 2021; Latchininsky et al., 2016; Steinkraus et al., 2017;

Wesołowski & Wesołowski, 2013). One of the largest known groups of parasites that cause

summit disease are the "zombie fungi" of the genus *Ophiocordyceps*, which infect various species

of insects and arachnids globally (Arruda et al., 2021; Cooley et al., 2018; de Bekker et al., 2021).

This includes wasps (e.g., *O. humberti* (Sobczak et al., 2020)), beetles (e.g., *O. curculionum*

(Shrestha et al., 2016)), scale bugs (e.g., *Ophiocordyceps clavulata* (Petch, 1933)), caterpillars

(e.g., *O. sinensis* (Guo et al., 2021)), flies (e.g., *O. dipterigena* (Chhetri et al., 2020)), and spiders

(e.g., *O. engleriana* (Sung et al., 2007)). However, a large portion of the currently described

species belong within the species complex *Ophiocordyceps unilateralis*; one of the more well-

studied groups of zombie fungi that collectively infect and manipulate carpenter ants with extreme host-specificity (Araújo et al., 2018, Araújo et al., 2020).

Ophiocordyceps unilateralis species begin their lifecycle as a spore, ejected from their parental fruiting body. These spores sail on the wind to infect new ants, either through immediate contact or through the production and attachment of secondary capillispores from the forest floor (Evans et al., 2018). Once inside the ant, the fungus begins to secrete an array of biomolecules in a time-specific manner; some of which alter the behaviour of their host (de Bekker et al., 2015; Will et al., 2020). The first notable change occurs as these ordinarily social hosts begin to show a lessened ability to communicate and participate in foraging tasks (Trinh et al., 2021). Subsequently, infected ants abandon the nest and climb nearby vegetation. Once at an elevated position, they latch on with their mandibles, such that the ant is perched on the underside of its biting substrate (Andersen et al., 2009). The fungus then completes its lifecycle through the formation of an endosclerotium that siphons the remaining nutrients from the host for the formation of a new fruiting body (Fredericksen et al., 2017). Closer observation and quantification of behaviours in both the lab and the field have led to the identification of other, more subtle behavioural changes. Altered behaviours include changes in daily rhythms (Hughes et al., 2011; Trinh et al., 2021) and the loss of motor coordination and balance, resulting in ants that fumble their way around as in a "drunkard's walk" (Supplemental Video 1) (Hughes et al., 2011; Trinh et al., 2021; Will et al., 2020). These conspicuous behavioural changes caused by the fungus during infection make the *O. unilateralis* species complex and their hosts particularly well-suited for the study of behaviour-manipulating parasites in greater detail (de Bekker et al., 2015).

Recent work establishing fungal culturing and ant infection techniques, as well as the genomic sequencing of several *O. unilateralis s.l.* species, has laid the groundwork for research

aimed at understanding the molecular framework driving behavioural manipulation (de Bekker et

al., 2015; de Bekker et al., 2017; Will et al., 2020). Transcriptomics analyses of the North

American species *Ophiocordyceps kimflemingiae* and *Ophiocordyceps camponoti-floridani* have

identified secreted proteins and secondary metabolite clusters highly upregulated during

summiting behaviour (de Bekker et al., 2015; Will et al., 2020). One of these secondary metabolite

clusters demonstrated a ~5,900 (in *O. kimflemingiae*) and ~12,000-fold (in *O. camponoti-floridani*)

increase in expression during manipulation compared to fungi grown in culture (Will et al., 2020).

This cluster was first identified as an ergot alkaloid-producing pathway based on the functional

annotation of its tryptophan dimethyltransferase (Pfam: TRP_DMAT (PF11991)) backbone gene

(de Bekker et al., 2015). More detailed annotation of the genes within and surrounding the cluster

demonstrated that it was homologous to the aflatrem-producing cluster in the plant pathogen

Aspergillus flavus (Will et al., 2020). Genomic comparisons with other *Ophiocordyceps* genomes

revealed that this secondary metabolite cluster appears to be conserved across species within *O.*

unilateralis but not in other *Ophiocordyceps* species outside the species complex (e.g.,

Ophiocordycpes australis s.l.) (de Bekker et al., 2017). This begs the question; what evolutionary

purpose may aflatrem-like products serve for the fitness of *Ophiocordyceps* species that

manipulate the behaviour of *Camponotus* hosts?

Aflatrem, a secondary metabolite of the indole-diterpene group called aflatoxins, is a potent

tremorgenic and carcinogenic mycotoxin known to cause neurological disorders in animals (Duran

et al., 2006; Nicholson et al., 2009). Cows exposed to aflatrem through ingestion of *A. flavus*-

contaminated grain feed exhibit neurological problems exhibited by muscle tremors and

hyperexcitability, followed shortly by incoordination, ataxia, and seizures (Selala et al., 2008;

Valdes et al., 1985). These neurological symptoms, linked directly to aflatrem's positive allosteric

modulation of Gamma-aminobutyric acid (GABA) receptors, are referred to as "staggers

syndrome" (Eldefrawi et al, 1990; Grant et al., 1987; Yao et al., 1989). Should aflatrem-like

compounds exhibit similar effects in *Camponotus* species, it could have important ramifications

for host manipulation given the suspected involvement of GABA receptors in caste and

behavioural differentiation in *Camponotus* ants (Gospocic et al., 2017; Graham, 2018). The

likelihood that an aflatrem-like compound may be employed during the manipulation of carpenter

ants is further supported by the "drunkard's walk" observed during late-stage infection; a

behaviour reminiscent of the staggers syndrome detailed in vertebrates (Supplemental Video 1).

We, therefore, hypothesize that aflatrem-like compounds are responsible for the stagger phenotype

observed during the behavioural manipulation of carpenter ants by *O. unilateralis* s.l.

In this study, we administered aflatrem to the Florida carpenter ant *Camponotus*

floridanus, the host of *O. camponoti-floridani*. We used microinjections to mimic the *in vivo*

secretion of aflatrem-like compounds by fungal cells in the haemolymph during infection (Figure

1).

Figure 1. Experimental overview of (A) the treatment groups, each consisting of 7 individuals from 3 different colonies of *Camponotus floridanus*, and an example of microinjection, (B) the arena setup for behavioural assays and parameters for the observational periods, and (C) the behavioural analyses using machine learning and behaviour scoring software.

Given the ease of its procurement, and its previous use in other animal models, we used high purity synthesized aflatrem (>98%) in our experiments (TePaske et al., 1992; Trienens & Rohlfs, 2011). Following injections, we observed and quantified ant behaviour to address if aflatrem can induce behaviours consistent with those exhibited during infection by *O. camponoti-floridani*. Furthermore, to investigate which host pathways are affected by aflatrem and potentially give rise to the exhibited behavioural phenotypes, we also conducted transcriptomics analyses of aflatrem-injected individuals. Comparison of these results to transcriptomics data from *Ophiocordyceps*-infected *C. floridanus* ants (Will et al., 2020) yielded several consistent

differentially expressed genes, including genes that encode neuromuscular and chemosensory

proteins.

2. MATERIALS AND METHODS

2.1 Ant Collection

For use in behavioural and transcriptomics experiments, we collected three separate colonies of

Camponotus floridanus using a DeWALT Cordless 2 Gallon Wet/Dry ShopVac at the University

of Central Florida's arboretum, coordinates: Colony 1 (28.602748, -81.194269), Colony 2

(28.603918, -81.191682), and Colony 3 (28.601430, -81.191251). We housed these colonies in 9.4

L plastic containers (Rubbermaid) lined with talcum (Acros Organics), applied using a viscous

20% w/v solution of powder suspended in 100% ethanol to prevent ants from climbing the walls.

Throughout the study, we provided the ants with fresh supplies of both double distilled water

(ddH2O) and 15% sucrose solutions, *ad libitum*. We also kept the housing in an incubation

chamber (Percival) set to the following program: 20°C – 0 lux – 80% relative humidity (RH) with

a linear increase in temperature, light, and decrease in humidity over 4 hours to reach 28°C – 2,100

lux – 65% RH, at which conditions were held for 4 hours followed by a linear decrease in the

temperature, light, and increase in humidity, returning to 20°C – 0 lux – 80% RH over 4 hours;

and held for 12 hours.

2.2 Microinjections

To test our hypothesis and identify the effect of aflatrem-like compounds on *C. floridanus*,

we established a behavioural assay using microinjections to introduce aflatrem in a manner

biologically relevant to the secretion of aflatrem-like compounds by *Ophiocordyceps* species in

the haemolymph of their ant hosts (Figure 1). For these assays, we utilized high purity aflatrem,

>98% $C_{32}H_{39}NO_4$, manufactured from *A. flavus* by BioCrick Science Solution Specialists in non-

water-soluble crystal form, which we suspended in 100% acetone (Fisher Chemical) to a

concentration of 10 mg/mL. We performed injections using borosilicate glass micropipettes

(Fisher) pulled with a glass puller (model: PC-100, Narishige) under the following parameters:

single step, all weights, heated to 62°C. To inject the ants, we held each one by the dorsal side of

the thorax before piercing the carapace on the ventral side between the first and second pair of

legs, followed by slowly injecting 0.5 µL of solution into the ant while simultaneously releasing

grip to alleviate intracavitary pressure.

Because the aflatrem used in this study is non-water soluble, we performed a preliminary

trial to identify acetone tolerance in ants. As such, we injected subjects with 100%, 50%, 25%,

10%, and 5% acetone solutions diluted with phosphate buffered saline (PBS) (137 mmol NaCl,

2.7 mmol KCl, 10 mmol Na_2HPO_4, 1.8 mmol KH_2PO_4, pH adjusted to 7.4), along with a 100%

PBS solution as a control. Ten minutes prior to injection, we removed containers housing ant

colonies from the incubator and set them on the lab bench top to acclimate to lab conditions. We

also performed all injections in triplicate, one ant per colony per treatment. After injection, we

placed the ants into individual 60 mm x 15 mm petri dishes (USA Scientific) and returned them to

the incubator. We kept these ants under observation for 48 hours post injection using an infrared

GoPro Hero6 set to record time-lapse footage with 60 second intervals and a CMVision IR30

Illuminator lamp for observation during the night cycle. From these videos, we were able to easily

differentiate dead and resting *C. floridanus* ants from one another; dying ants coiled their legs and

fell to one side upon death while resting ants remained prone with legs supporting their bodies. As

such, we were able to estimate the time of death of each ant within an interval of 1 minute by

noting the frame in which they exhibited "death behaviours". We euthanized any ants that survived

the 48-hour test period by freezing at -20°C for 24 hours. Results from these preliminary analyses

indicated that a 10% acetone solution was the highest concentration for which there were no

statistical differences in average survivability compared to PBS-injected and non-injected controls.

To test the survivability of ants post injection, and determine if any observed behavioural

phenotypes may be attributed to "dying behaviours" resulting from aflatrem exposure, we

performed a second preliminary trial. We injected 7 ants with 0.5 μL 10% acetone solution

containing 500 ng aflatrem. This was the highest dose possible using the 10 mg/mL aflatrem stock.

After injection, we recorded the ants for 48 h using the same GoPro setup as the previous

preliminary test. We then compared the two groups and found no significant differences in

mortality, with all individuals surviving for at least 24 h and an average time of death arising at 34

h. The results indicate that all ants that died in the trial did so as a result of desiccation from lack

of water.

Confident that neither 10% acetone nor 500 ng aflatrem killed the ants or induced death-

related behaviours, we conducted an assay to determine the effects of aflatrem on ant behaviour

using 10% acetone solutions containing the following doses of aflatrem; 10 ng, 50 ng, 100 ng, 500

ng. For each dose of aflatrem, we injected 7 ants from each of the 3 colonies (N=21, per dose) for

a total sample size of N=126. We individually lifted each ant from their housing by hand wearing

neoprene gloves (NeoTouch) and allowed them to acclimate to the lab conditions by placing them

in individual 60 mm x 15 mm petri dishes (USA Scientific) on the lab benchtop for 10 minutes.

We fitted each petri dish with a thread of twine spanning the dish from the bottom of one side to

the top of the other. This provided ants with a substrate for biting and more dexterous climbing

behaviours (Figure 1-B). To allow ants to walk around the perimeter of the dish unimpeded, we

made sure to tape (Scotch) the twine pressed flat against the bottom corner of the petri dishes. We

created a new petri dish arena for each test subject (N=126) to avoid any behavioural changes that

may be induced via chemical ques left by prior occupants. After the 10-minute acclimation period,

we injected ants with aflatrem following the microinjection protocols established in the

preliminary trials. The doses of aflatrem used were based on approximate ant equivalent weights,

verified by measuring *C. floridanus* body volume through length and weight following each

experiment. While we measured body lengths to the nearest 0.5 mm for each individual, weight

was measured in groups of seven ants corresponding to their treatment and colony ID. We recorded

these measurements immediately following euthanization at -20°C, preserving the approximate

water weight for each ant. To account for the approximate weight of solution injected into each

ant, we also subtracted 3.5 mg from the total weight of each group. Based on an average of 8.28

mg weight and 6.68 mm length calculated from measurements of all 126 subjects used in this study

(Supplemental Table 1), the four doses of aflatrem that were used are approximately 6.04e-5,

1.21e-5, 0.60e-5, and 0.12e-5 ant equivalents. These ant equivalent doses are consistent with

pervious works testing insecticidal compounds in ants (Buczkowski & Woosler, 2019; Klotz &

Moss, 1996; Sakamoto & Goka, 2021).

2.3 Video scoring of behavioural assays post injection

To record behaviours post injections, we used an iPhone 13 pro. We recorded three or four arenas

at a time in a rolling format to ensure high quality resolution at 60 fps. This generated video files

approximately 45 minutes in length. In cases where ants exhibited signs of physical injury from

the injection, e.g., immobility in the hind legs or body curling, we euthanized the subject and

replaced it with a new ant and new arena. If any ants were exhibiting resting behaviour at the end

of their recording period, we tapped their arenas several times to induce movement. All instances

of tapping elicited activity, confirming that all test subjects were still alive.

To crop the rolling videos into clips showing only individual arenas, we first converted the

iPhone .MOV files to .mp4 files using iMovie. We then exported these .mp4 files to Movavi Video

Editor Plus 2022 and trimmed each clip to exactly 30 minutes, beginning as soon as each arena

was positioned in the field of view. The names for each of these clips were blinded before two

observers manually scored the videos using the behaviour tracking software CowLog3 (Pastell,

2016). Before formal scoring, we trained each observer using eight test run videos; four of ants

injected with 500 ng aflatrem, and four of ants injected with PBS. These videos were made during

proof-of-principle testing and are not included in the data presented in this study. We used the

results from these practice runs to form a consensus on scoring methodology. From these training

videos, the observers identified eight scorable behaviours: Climbing Wall, Climbing Top,

Climbing Rope, Walking, Staggering, Grooming, Resting, and Biting (Supplemental Video 2).

While most of these behaviours are self-explanatory, we defined "staggering" as a slow, sluggish

walking behaviour with repeated lateral movements (Supplemental Video 3). We also categorized

behaviours as "staggering" when trouble balancing, listing to one side, falling over, walking in

tight circles, flipping over, or dramatic wagging of the gaster were observed (Supplemental Video

3). Additionally, each observer kept notes on unique behaviours observed during the scoring

period, including walking backwards, lunging/bucking, and two instances of convulsions

(Supplemental Video 4).

During the formal scoring, we assigned the blinded videos to the observers in a randomized

block format (Supplemental Figure 1-A) with each observer scoring three videos at random per

treatment group per colony. Both observers scored the seventh video for each treatment group,

from which the results were compared and used as a formative metric to ensure consistent scoring

throughout the evaluation period. While most behaviours were readily distinguishable, resulting

in very similar scores between observers, walking and staggering provided the most discrepancy.

We, therefore, used staggering as the primary measurement to evaluate scoring consistency across

observers (Supplemental Figure 1-B).

In addition to manual observations, we also analysed each video using the MATLAB-based

machine learning tool Massively Automated Real-time GUI for Object-tracking (MARGO)

(Werkhoven et al., 2019). Using this tool, we were able to measure the speed and distance travelled

by each subject. While we calculated distance travelled by measuring the total distance between

centroid coordinates over the entire 30 min duration, we only measured speed, calculated in mms

travelled per second, during activity. To optimize the tracking within the field of view for each

video, we used the following metrics: region of interest (ROI) was set to circular shape and

manually overlayed around the arena, background referencing was set to detect dark objects on

light backgrounds, tracking threshold was set between 70-80, distance of the arena was set to 60

mm using the custom measuring tool, frame rate was set to 30, and min/max area was set to 5 and

80, respectively.

Before testing the effects of aflatrem on ant behaviour, we compared the results from the

10% acetone controls to the 100% PBS controls to identify any potential behavioural phenotypes

induced by the addition of acetone to the injections. To test for any statistical significance between

the two groups, we used a two-sample t-test in RStudio v1.3.959. Following the comparison of

these two controls, we tested the effect of aflatrem exposure on ant behaviour with a linear

regression model using the lme4 package in RStudio, by comparing the 10% acetone control (0 ng

aflatrem) with 10% acetone containing 10 ng, 50 ng, 100 ng, and 500 ng aflatrem treatments.

2.4 RNA sequencing

To understand the effects of aflatrem exposure in *C. floridanus* at the genomic level, we injected

a total of 10 ants with either 10% acetone (control group, N=5) or 500 ng aflatrem (treatment

group, N=5). We then placed each ant into their own petri dish arena for 4 minutes. We chose this

timepoint based on the scaling levels of activity observed in our behavioural assays (Supplemental

Figure 2). After this period, we quickly placed the ants into 2 mL microcentrifuge tubes and snap

froze them in liquid nitrogen. Subsequently, we removed the head from each ant with forceps and

placed it in a pre-frozen 2.0 mL Self-Standing Impact Resistance Microtube (USA Scientific)

containing 2 Grade 25 5/32" metal ball bearings (Wheels Manufacturing) for mechanical tissue

disruption in a SPEX® SamplePrep 1600 MiniG at 1500 rpm for 30 sec. We dissolved the frozen

samples in 500 uL TRIzol (Ambion) to extract the RNA and isolated it using the Qiagen RNeasy®

MiniElute® Cleanup Kit (Will et al., 2020). The resulting RNA was stored at -80C and shipped to

Azenta US for Next Generation Sequencing. Libraries were prepared by Azenta US using PolyA

selection and sequenced as 150 bp paired end reads on an Illumina HiSeq, resulting in ~350M

reads. Read quality was verified using FastQC (Andrews 2010). We used Trimmomatic (Bolger

et al., 2014) to trim reads for adapters and quality using the following parameters: Leading: 3,

Trailing: 3, MinLen: 36. We then aligned the transcripts to the *Camponotus floridanus* reference

genome (Gene bank ascension number GCA_003227725.1) using Spliced Transcripts Alignment

to a Reference (STAR) 2.7.10a (Dobin et al., 2013). After alignment, we checked the resulting

.bam files for quality using the standard metric of uniquely mapped reads > 70%, sorted, and

indexed using Samtools (Danecek et al., 2021). We next calculated the number of gene hits for

each sample using FeatureCounts, following the suggested counting of strictly uniquely mapped

reads within exonic regions (Liao et al., 2019). The quality of the resulting files was validated

using a benchmark of 15-20 million reads per sample. After removing hits with low level of

expression (counts less than 100 per gene, i.e., an average less than 10), we normalized the

remaining counts using DESeq2 (Love et al., 2014) eliminating 3,227 genes from further analysis.

We performed differential expression analysis on the remaining 10,443 genes using DESeq2 with

an alpha value of 0.05 and functionally annotated the differentially expressed genes using

Blast2GO (Conesa et al., 2005).

2.5 Ethical Note

We performed all injections and behavioural analyses in this study on the invertebrate ant species

Camponotus floridanus, which is not endangered and is a prevalent species in Central Florida. We

collected colonies of this species at the University of Central Florida Arboretum, which did not

require prior approval from an animal welfare or ethics committee. Only three colonies were

collected to meet the minimal scientific standard of triplicate replication. Collection utilized a fast

and handling-minimizing method. We applied the following standards in effort to reduce the

amount of stress incurred by the ants throughout our study as much as possible. Housing was

conducted in an environmental chamber set to mirror the subjects' natural conditions. We

maintained colonies by providing *ad libitum* food and water, and dark enclosures for nesting. We

furthermore conducted experiments on the minimum required number of subjects to reach the

sample size needed for statistical analysis. Ants were quickly euthanized at -20°C in instances

where they were injured during treatment and immediately following all experiments.

3. RESULTS

3.1 The behavioural effects of aflatrem in *C. floridanus*

Before we could analyse the behavioural effects of aflatrem in ants, we screened for any potential behavioural changes caused by the presence of 10% acetone in the injections. Using an independent two sample t-test to compare the behaviour of ants injected with PBS to those injected with 10% acetone, we were able to rule out any significant differences between the two groups: rest ($p = 0.787$), speed ($p = 0.997$), distance ($p = 0.947$), stagger ($p = 0.205$). Even though neither group was exposed to aflatrem, we unexpectedly observed a few instances of staggering. To determine if a small amount of staggering behaviour is normal in ants, which would need to be accounted for in our results, we reviewed the footage at timestamps logged for these events with an increased level of scrutiny. Upon closer review, we discerned that nearly all staggering observed in both groups fell into the "gaster wagging" sub-type (Supplemental Video 2), and nearly always occurred in the first 30 seconds of the recording. This trend was also observed in the aflatrem treatment groups as well, indicating that this particular type of staggering behaviour is likely a transient physical response to the trauma resulting from the injection process. We confirmed this hypothesis through a quick follow-up experiment pricking ants with empty glass capillaries, which also induced gaster wagging in some.

Confident that behaviours were not significantly affected by the addition of 10% acetone in the injections, we made comparisons across increasing doses of aflatrem. A linear regression model showed that time spent resting in the arena was positively correlated with increasing doses of aflatrem ($F_{4,105} = 5.54$, $p = 2.05e-2$) (Figure 2-A), while the average speed of aflatrem-treated ants was negatively correlated ($F_{4,105} = 4.91$, $p = 2.89e-2$) (Figure 2-B). However, the average

distance travelled did not correlate with aflatrem treatment ($F_{4,105}$ = 3.036, p = 8.44e-2, ns) (Figure

2-C). Finally, bouts of staggering did positively correlate with increasing doses of aflatrem ($F_{4,105}$

= 36.65, p = 2.32e-8) (Figure 2-D).

Figure 2. Behavioural results collected from *C. floridanus* ants injected with increasing doses of

aflatrem. Each boxplot depicts the quartiles of the distribution for each treatment with ants from

colonies 1, 2, and 3 shown as purple, light blue, and dark blue dots, respectively. A) shows the

proportion of time spent resting for each ant, B) shows their average speed during activity in

millimetres per second, C) shows their total distance travelled in meters, and D) shows the

proportion of activity spent staggering for each ant. The linear regression model results are shown

in the centre of the graph for each behaviour. A & D) both contain pie charts illustrating the relative

duration of each behaviour, while B) contains speedometers for each treatment and C) presents the

median tracer plot from MARGO for each group.

3.2 The effects of aflatrem on gene expression in *C. floridanus*

While injection of *C. floridanus* with aflatrem resulted in quantifiable phenotypes, we next asked

which genes could be underlying the emergence of staggering and diminished activity phenotypes.

As such, we conducted transcriptomics to compare gene expression in the heads of ants treated

with 500 ng aflatrem (treatment group) to that of ants treated with 10% acetone solutions (control

group). We found 261 genes that were significantly differentially expressed (Supplementary File

1). Of these genes, 148 were upregulated and 114 were downregulated in the treatment group

(Figure 3). We found the largest increase of expression (~22-fold increase) in a gene encoding an

aminopeptidase N-like protein, a common cell surface hydrolase involved in a variety of cellular

processes (Nocek et al., 2008). The largest decrease in expression (~512-fold decrease) was

exhibited by a gene coding for a cytochrome P450 6k1, a protein involved in detoxification of

insect cells (Xing et al., 2021) (Figure 3).

Figure 3. Volcano plot for the differential expression of 10,443 *Camponotus floridanus* genes in ants treated with 500 ng aflatrem compared to the 10% acetone control group. Dots in red represent non-significant changes in gene expression, while dots in blue represent significant changes using an alpha value of 0.05. Functional annotations are shown for genes with adjusted *P* values less than 5e-5 and absolute log$_2$ fold changes lower than -6 or greater than 3.5.

Several of the 261 differentially expressed genes were predicted to play a role in neuromuscular systems. Among them, we found an *anoctamin-4*, which codes for an ion channel protein involved in Ca^{2+}-dependent conductance (Reichhart et al., 2019), downregulated in

aflatrem-treated ants. We also detected a *membrane metallo-endopeptidase* (*Neprilysin*), a

transmembrane protein found in neurological tissue that plays a role in tissue perception and

preservation (Obulesu, 2019), which was slightly upregulated in aflatrem-treated ants. More

specific to muscular function, we found 2 upregulated *titins* genes, which code for very large

proteins that play an essential role in sarcomere function (Freiburg et al., 2000; Zhang et al., 2000).

Also upregulated was a *ryanodine receptor* gene, responsible for releasing Ca^{2+} in the smooth

endoplasmic reticulum of muscle cells during excitation-contraction coupling (Lanner et al., 2010;

Shakiryanova et al., 2007). On the other hand, *caveolin-3*, which codes for a scaffold protein that

organizes signalling molecules in the membranes surrounding muscle cells (Dewulf et al., 2019;

Galbiati et al., 2000-2013) was found to be downregulated, as well as 2 *myosin ID heavy chain-*

like protein genes that act as the motor proteins of muscles (Wells et al., 1996) (Supplementary

File 1).

In addition to neuromuscular genes, we also identified many downregulated olfactory-

related genes, including 4 general odorant-binding protein genes (OBPs) (i.e., *OBP19a*, *OBP56a*,

and 2 *OBP69a*). These proteins are located in the antennal sensillary lymph of insects and bind to

pheromones that play a role in insect communication and interactions (Gaubert et al., 2020).

Moreover, we found 3 significantly downregulated *ejaculatory bulb-specific* genes, involved in

the release of pheromones in insects (Pelosi et al., 2018), and a downregulated *sensory neuron*

membrane protein gene, which has been found to play a role in pheromone sensitivity in the moth

Heliothis virescens (Pregitzer et al., 2014).

We also detected a differentially expressed gene coding for a circadian clock-controlled

protein. Such genes could be of interest, given previous explorations of the time-dependent nature

of behavioural manipulation and the disruption of clock-dependent foraging behaviours in

Ophiocordyceps-infected ants (de Bekker et al., 2015; de Bekker & Das, 2022; Will et al., 2020),

This gene, which showed an 18-fold decrease in expression in aflatrem-injected ants compared to

10% acetone controls, also contains a Haemolymph Juvenile Hormone Binding Protein Pfam

domain. Such binding proteins, driven by circadian rhythms, are known to act as carrier proteins,

chaperoning juvenile hormones, which play a wide array of roles in insect physiology and

development, to their target tissues (Zalewska et. Al., 2009).

To investigate if any of the differentially expressed genes were affected during

Ophiocordyceps-induced behavioural manipulation, we compared our RNA-seq results to

transcriptomics data obtained from *C. floridanus* manipulated by *O. camponoti-floridani* (Will et

al., 2020). We found that 108 of the 261 differentially expressed genes observed in aflatrem-treated

ants were also differentially regulated in late-stage *O. camponoti-floridani* infections

(Supplementary File 1). Of these genes, 34 were differentially regulated in the same direction,

while 74 shared an inverse relationship. The previously mentioned *cytochrome P450*, *sensory*

neuron membrane protein encoding gene, *OBP56a*, and *caveolin-3* genes were amongst the

similarly expressed genes, all of which were downregulated. We also found an upregulated *nuclear*

hormone receptor E75, member of a family of transcription factors that are known to be activated

by juvenile hormone signalling pathways in Diptera and Lepidoptera (Dubrovskaya et al., 2004).

In addition, we detected an upregulated circadian clock steroid-producing gene linked to the

maintenance of circadian rhythms during stress in *Drosophila* (Kumar S et al., 2014). Moreover,

a *neurogenic locus protein delta*, capable of coding for multiple translational products including

neural differentiation factors and growth hormones (Kopczynski et al., 1988), was also upregulated

in both datasets.

Amongst the differently expressed genes, the aforementioned *OBP19a-like* gene, 5

cytochrome P450s, and an *ejaculatory bulb-specific protein 3*, were all significantly

downregulated in aflatrem-treated ants but upregulated in *Ophiocordyceps*-infected ants. More

genes shared this same inverse relationship, including an *odorant receptor coreceptor* gene, coding

for proteins expressed in chemosensory organs that form heteromultimeric complexes within

odorant receptors (Mukunda et al., 2014). We also detected a *short neuropeptide F*, which is

implicated in the regulation of animal behaviour including circadian rhythms and hunger (Lee et

al., 2004), and a *netrin receptor UNC5C*, which aids in the formation of axon connections through

axon extension and cell migration (Kim & Ackerman, 2011). S*ialin*, a sialic acid transporter-

coding gene that prevents neurodegenerative disorders caused by accumulation of sialic acid

(Tarailo-Graovac et al., 2017) was additionally among these differentially expressed genes.

Upregulated in our aflatrem-treated ants, but downregulated in the *Ophiocordyceps*-

infected ants, was a *receptor-type tyrosine-protein phosphatase delta* (*PTPRD*). This gene is part

of the PTP family involved in many known signalling pathways for which delta is shown to

promote neurite growth and regulate axon guidance (Tomita et al., 2020). In addition, we found

an *obscurin*, which, like *titin*, codes for a large protein expressed in skeletal muscles that anchors

myofibrils to the sarcoplasmic reticulum for muscle contraction (Feher, 2017), and a *juvenile*

hormone acid O-methyltransferase-like gene, which codes for an enzyme that converts precursor

molecules into juvenile hormones (Shinoda & Itoyama, 2003). Finally, the two data sets also

shared 26 genes with statistically significant changes in expression coding for proteins of unknown

function (Supplemental File 1).

4. DISCUSSION

Certain parasites can induce host behaviours that increase the reproduction, survivability, and

transmission of those parasites at a cost to the host. How such behavioural manipulations are

established is still poorly understood. In recent years, multi-omics work on "zombie ants" has

provided a myriad of candidate compounds that *Ophiocordyceps* fungi might use to manipulate

ant behaviour. However, their causal relationship to resulting host behaviours still needs to be

functionally tested. This study aimed to test aflatrem-like compounds previously implicated in the

establishment of *Ophiocordyceps* extended phenotypes in zombie ants (Will et al., 2020). We

accomplished this by exposing healthy *C. floridanus* ants to high purity aflatrem and quantifying

its behavioural effects. Following injections, we found dose of aflatrem to be positively correlated

to the amount of time that treated individuals spent staggering. To rule out that this observed

behaviour was a symptom related to death, we performed a longevity assay to demonstrate that

aflatrem does not kill *C. floridanus*. These preliminary tests demonstrated that aflatrem-like

compounds are unlikely to be employed by *Ophiocordyceps* as a means to kill hosts. Instead, the

observed staggering behaviours indicate that the aflatrem-like compounds produced by

Ophiocordyceps during infection are, at least in part, responsible for the drunkard's walk

phenotype exhibited by infected carpenter ants (Hughes et al., 2011; Trinh et al., 2021). How the

emergence of this behaviour would adaptively contribute to *Ophiocordyceps* fitness is, however,

unknown.

Our experiments further demonstrated that aflatrem has a significant effect on the mobility

of ants, decreasing overall activity as dose increased. Ants injected with higher doses of aflatrem

moved less often, did so much slower when they were active, and as a result travelled shorter

distances. We also observed that they had more difficulties climbing. Particularly when it involved

the twine in the arena, which requires greater dexterity than climbing on the wall or top of the petri

dish. Previous field studies suggest that *Ophiocordyceps* development and fruiting body formation,

needed for spore transmission, are dependent on precise levels of light, humidity and temperature

(Andersen & Hughes, 2012; Andriolli et al., 2019; Cardoso Neto et al., 2019; Hughes et al., 2011;

Lavery et al., 2021; Will et al., under review). Aflatrem-mediated tremors and reduction of

dexterity and speed could aid to keep the ant within these optimal microclimates prior to biting,

by preventing it from climbing too high into the canopy.

While we could reason a parasite-adaptive function for aflatrem-reduced activity, it

contradicts previous studies where *Ophiocordyceps*-infected individuals showed hyperactive

locomotion prior to summiting (de Bekker et al., 2015; Trinh et al., 2021; Will et al., 2020).

Nevertheless, transcriptomics studies on *Ophiocordyceps*-manipulated ants suggest that the fungus

expresses manipulation compounds in a time-specific manner (de Bekker et al., 2015; Will et al.,

2020). It is reasonable to suggest that, prior to the upregulation of aflatrem-like compounds,

Ophiocordyceps secretes effectors that induce hyperactivity and climbing behaviours to drive the

ant away from the nest and up nearby structures. An example of one such effector could be protein

tyrosine phosphatase (PTP). Genes encoding this protein are upregulated during *Ophiocordyceps*

manipulation (de Bekker et al., 2015; Will et al., 2020) and PTPs have been found to induce

hyperactivity and enhanced locomotion in caterpillars that are infected with a behaviour-

manipulating baculovirus (Han et al., 2015). Enhanced locomotion could be parasite-adaptive by

leading hosts away from aggressive conspecifics, which can recognize infected individuals and

attack them (Trinh et al., 2021; Will et al., 2020) as part of their social immunity (Cremer et al.,

2007). After the ant is safely away from the adaptive behaviours of the colony, the production of

debilitating compounds could function to maintain the ant in a desirable, elevated location.

In addition, aflatrem-like compounds may also aid in the positioning of the host for the

final, and perhaps most iconic, behavioural manipulation; the "death grip". During the death grip,

ants bite onto vegetation with their mandibles, firmly anchoring them in place. Shortly after, the

fungus kills the host and quickly consumes the remaining nutrients to produce its reproductive

structures. The attachment of the host is a vital step in host manipulation as it promotes infective

spore formation and release. Without it, ants would fall back to the forest floor, which interferes

with fruiting body formation and, eventually, transmission (Andersen et al., 2012; Andriolli et al.,

2019; Loreto et al., 2014; Will et al., under revision). The large increase in resting behaviours seen

in ants that received the highest dose of aflatrem could aid in positioning of the ant's head for

biting behaviours. Moreover, aflatrem injection resulted in the differential expression of muscle-

related genes in the heads of treated ants as compared to the control group. Therefore, the muscle-

related effects of aflatrem-like compounds secreted by *Ophiocordyceps* could potentially be

directly involved in the locking of the mandible muscles as part of the biting phenotype (Hughes

et al., 2011). However, we did not observe increased biting behaviour in aflatrem-treated ants in

this study.

Figure 4. Examples of "hugging" behaviour portrayed by *Ophiocordyceps*-infected *Camponotus*

floridanus ants during the death grip observed in a natural setting.

As part of the death grip phenotype, *O. camponoti-floridani*-infected *C. floridanus* are

often found wrapping their limbs around thin pieces of Spanish moss, which they most frequently

use as a biting substrate (Figure 4) (Will et al., under review). This "hugging" behaviour has also

been observed in other *Ophiocordyceps*-ant interactions where smooth substrates, such as twigs,

are primarily bitten (Loreto et al., 2018). The transcription-level dysregulation of sensory

perception and neuromuscular functioning that we found, as well as the observed staggers, suggest

that unsteadiness during climbing might cause ants to stumble and hug the vegetation. As such,

the hugging behaviour itself could act as a secondary means by which hosts are kept in elevated

positions should biting itself fail. Indeed, the adherence to vegetation through biting is sometimes

missing in *O. camponoti-floridani*-manipulated ants, both in the field and in the lab (Will et al.,

2020) while hugging is always observed. In fact, our group has witnessed multiple examples of

such instances in the field where hosts with fully grown fruiting bodies are held to strands of

Spanish moss by just their limbs, either due to failed biting or the complete loss of the ant's head

from the sample. Hugging may be unique to ant hosts of *Ophiocordyceps* species as the phenotype

is absent from other host-manipulating entomopathogens, such as *Eryniopsis lampyridarum*

infecting the Goldenrod Soldier Beetle, which similarly induces summiting and death grip

behaviours on thin, huggable foliage (Steinkraus et al., 2017).

The exact structure and concentrations of the aflatrem-like compounds produced by

Ophiocordyceps in vivo are still unknown. Regardless, we were able to use reasonable doses of

purified aflatrem (i.e., within body weight equivalents used in previous pesticide testing) to

replicate the staggers/"drunkard's walk" phenotype induced in ants during *Ophiocordyceps* infection. This caused the differential expression of 108 ant host genes that were previously identified in *Ophiocordyceps*-infected ants exhibiting the summiting phenotype. As such, at least in part, similar host pathways appear to be targeted during isolated aflatrem-dosing and *Ophiocordyceps* infection, leading to comparable behaviours. This indicates that aflatrem-like compound secreted by *Ophiocordyceps* is likely responsible for some of the transcriptional and behavioural changes observed during manipulation. Future work characterizing the chemical composition and three-dimensional structure of aflatrem-like compounds produced by species of the *O. unilateralis* complex is needed to shed further light on the physiological effects of these alkaloids and their role in behaviour manipulation. Nevertheless, our results demonstrate that exposing hosts to hypothesized parasite manipulation compounds is a valuable avenue to test their proposed effects at the genome and phenome level. As such, integrative behavioural studies like this one can bring us one step closer to characterizing the molecular mechanisms that drive zombie-making fungi. Furthermore, the protocols established as part of this research provide a framework for future studies that wish to examine the roles of putative effectors produced and secreted by *Ophiocordyceps* species, paving the way for further characterization of host-manipulating compounds.

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Author Contributions:

WCB and CdB designed the experiment, analysed the data, performed the RNA extractions, and

wrote the paper. WCB, CK, and UAM collected the ants, performed the injections, and recorded

the video footage. CK and UAM performed the behavioural analyses using CowLog. WCB ran

the MARGO software and performed the statistical analyses.

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Supplemental Files

Supplemental Table 1: The average length and weight of *Camponotus floridanus* in this study

Colony #	Treatment Group	Average Length (mm)	Average Weight (mg)
Colony 1	C1-PBS	7.07	7.56
	C1-PBS/10% acetone	6.50	6.93
	C1-PBS/10% acetone/500ng aflatrem	6.78	7.94
	C1-PBS/10% acetone/100ng aflatrem	7.00	8.14
	C1-PBS/10% acetone/50ng aflatrem	6.43	8.44
	C1-PBS/10% acetone/10ng aflatrem	7.29	9.01
Colony 2	C2-PBS	7.07	9.49
	C2-PBS/10% acetone	7.07	10.27
	C2-PBS/10% acetone/500ng aflatrem	6.93	11.09
	C2-PBS/10% acetone/100ng aflatrem	6.93	10.48
	C2-PBS/10% acetone/50ng aflatrem	6.86	10.86
	C2-PBS/10% acetone/10ng aflatrem	6.86	10.13
Colony 3	C3-PBS	6.07	5.40
	C3-PBS/10% acetone	6.93	7.93
	C3-PBS/10% acetone/500ng aflatrem	6.79	6.44
	C3-PBS/10% acetone/100ng aflatrem	5.71	5.40
	C3-PBS/10% acetone/50ng aflatrem	5.64	6.23
	C3-PBS/10% acetone/10ng aflatrem	6.29	7.24
	Total	6.68 mm	8.28 mg

Supplemental Figure 1

Supplemental Figure 1. An overview of the randomized block format used to assign videos to each

observer (A). The videos assigned to observer 1 are shown in green, the videos assigned to observer

2 are shown in blue, and the videos assigned to both for formative analysis are shown in grey. B)

depicts the duration of staggering scores for each grey assignment, with the scores for observer 1

shown as green squares and the scores for observer 2 shown as blue circles.

Supplemental Figure 2

Supplemental Figure 2. Bar graphs depict the number of ants exhibiting staggering behaviour

(black) and the number of ants resting (dark grey) across the 30-minute observational period,

broken down into 5 second intervals. Control refers to ants injected with 10% acetone solvent.

Low dose aflatrem shows the data for ants that received 10-50 ng aflatrem, while high dose

aflatrem visualizes staggering and resting for ants that received 100-500 ng.

Supplemental Figure 3

Supplemental Figure 3. The path travelled by each ant recorded by MARGO tracking software

during the 30-minute observational period is shown in dark blue. The three columns on the left

show the paths travelled by the control group injected with 10% acetone, while the three columns

on the right show the paths travelled by ants in the 500 ng aflatrem treatment group. Each group

is further divided by colony ID using purple, blue, and dark blue outline to indicate individuals

from colonies 1, 2, and 3, respectively.