Glyphosate Inhibits Melanization and Increases Susceptibility to Infection in Insects

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ABSTRACT

Melanin, a black-brown pigment found throughout all kingdoms of life, has diverse biological functions including: UV protection, thermoregulation, oxidant scavenging, arthropod immunity, and microbial virulence. Given melanin's broad roles in the biosphere, particularly in insect immune defenses, it is important to understand how exposure to ubiquitous environmental contaminants affects melanization. Glyphosate – the most widely used herbicide globally – inhibits melanin production, which could have wide-ranging implications in the health of many organisms, including insects. Here, we demonstrate that glyphosate has deleterious effects on insect health in two evolutionary distant species, *Galleria mellonella* (Lepidoptera: Pyralidae) and *Anopheles gambiae* (Diptera: Culicidae), suggesting a broad effect in insects. Glyphosate reduced survival of *G. mellonella* caterpillars following infection with the fungus *Cryptococcus neoformans* and decreased the size of melanized nodules formed in hemolymph, which normally help eliminate infection. Glyphosate also increased the burden of the malaria-causing parasite *Plasmodium falciparum* in *A. gambiae* mosquitoes, altered uninfected mosquito survival, and perturbed the microbial composition of adult mosquito midguts. Our results show that glyphosate's mechanism of melanin inhibition involves antioxidant synergy and disruption of the reaction oxidation-reduction balance Overall, these findings suggest that glyphosate's environmental accumulation could render insects more susceptible to microbial pathogens due to melanin inhibition, immune impairment, and perturbations in microbiota composition, potentially contributing to declines in insect populations.

Key Words: melanin, phenoloxidase, malaria, midgut microbiome, microbiota *Galleria mellonella*, *Anopheles,* vector biology, nodulation, fungi, tyrosinase, hormesis, antioxidant, acid synergism

INTRODUCTION

Melanin, a black-brown pigment found in all biological kingdoms, is produced through a series of oxidation and reduction reactions. These reactions are typically catalyzed by two classes of enzymes: laccases (EC. 1.10.3.2) and phenoloxidases – a family which includes tyrosinases (EC. 1.14.18.1) (1). Tyrosinases are copper metalloenzymes found in bacteria, fungi, protists, arthropods, birds, and mammals (2–7), and have two catalytic roles: 1) hydroxylation of monophenols into *ortho*-diphenols, followed by 2) oxidation of *o*-catechols into *o*-quinones (8). During melanization, tyrosinase converts 3,4-dihydroxyphenylalanine (L-DOPA) to dopaquinone. Dopaquinone undergoes oxidation and reduction reactions to first form dopachrome, then dihydroxyindole. Dihydroxyindole undergoes radical-mediated polymerization to form melanins (8,9).

Melanization is an important component of immunity in virtually all insects (9). Upon infection, protease cascades are activated that cleave pro-phenoloxidases into active phenoloxidases. Phenoloxidases convert catecholamines in the hemolymph into melanin, which surrounds and eliminates the pathogen through exposure to reactive oxygen species (ROS) and lysis from toxic melanin intermediates (3,9–13). This melanization process is a key component of insect immune defense against bacterial, fungal, and protozoan pathogens, nematode parasites, and insect parasitoids (14–21). In addition, pathogens are cleared by two similar processes: nodulation of smaller microbes such as bacteria, fungi, and protozoa, and encapsulation for infections with larger organisms such as helminths and parasitoid eggs (22). Both nodulation and encapsulation involve pathogen neutralization via melanin accumulation and hemocyte (insect "blood cells") aggregation around the pathogen (22). Melanization and phenoloxidases are also important for wound healing and cuticular development – processes vital for insect health and survival (23,24). Since melanization is an essential physiological process and effector of insect health, understanding how common environmental contaminants affect melanin production is important. The significance of this is also highlighted by findings

suggesting that insect populations may be in decline in recent decades (25).

One ubiquitous chemical found in the environment is glyphosate, the most commonly

used herbicide world-wide, which was previously shown to interfere with melanization in the

fungus *Cryptococcus neoformans* (26). Glyphosate is a phosphonic glycine analogue and the

active ingredient in Roundup herbicides (27). It kills plants through competitive inhibition of

EPSP synthase in the shikimate pathway responsible for aromatic amino acid synthesis in many

plants, fungi, and bacteria (28). The low cost of glyphosate and wide availability of genetically-

modified glyphosate-resistant crops has increased both crop yields and glyphosate-based

herbicide use in agriculture (29,30). Between 1996-2014, glyphosate-resistant crops were linked

to a 12-fold global increase in glyphosate use, including 8-fold in the US, 134-fold in Brazil, and

107-fold in Argentina (31,32).

In practice, glyphosate is commonly applied at concentrations of ~28 to 57 mM (33) or in

formulations of 360 g/L (2 M), with 720 g (4 mol) per hectare (34). Glyphosate-based herbicides

are sprayed onto crops where the glyphosate is taken up by plant leaves and translocated to

growing tissues throughout the plant (35). Glyphosate is translocated to the roots where it is

released into the soil (34). In total, about 88% of the sprayed glyphosate ends up in the topsoil

(36–38)(38) . Less than 1% of glyphosate has been shown to enter water bodies, typically

following heavy rain, snowmelt, ploughing, or erosion (37), but concentrations from <1 nM to

~30 µM in nearby water have been reported (39). Further, glyphosate has been shown to enter

the air through wind erosion and deposit via rain (40).

Glyphosate is remarkably stable, with half-life ranging from weeks to years depending

on the surrounding microbial populations, which provide the primary mechanism of glyphosate

degradation, while temperature, light, acidity, and salinity also play roles in the degradation

process. Microbes mostly break down glyphosate into aminomethylphosphonic acid (AMPA),

which persists up to 20 times longer than glyphosate, and is often found in higher

concentrations in topsoil and water (41–45).

While glyphosate may have harmful effects on microbes and animals (as reviewed in

(35,46)), its impact on environmental microbial communities is inconclusive. Some studies

demonstrate clear perturbation of microbial communities, including disrupting rhizosphere

composition and fungal endophyte growth and viability (47–50), while others show little to no

long term impact on microbial communities (51–53), with no effects on overall soil health or

reduction in soil microbial mass (54). Microbial communities are also abundant in insect guts,

where they are important for insect health (55–58), and several studies have linked detrimental

effects of glyphosate on insect health to disruption of the microbiota. Honeybees exposed to

glyphosate have altered microbiomes and are more susceptible to *Serratia marcescens* (59),

although AMPA did not have the same effect (60). In tsetse fly midguts, glyphosate disrupts

Wigglesworthia glossinidia's production of folate - a compound required for tsetse fly health and

vector competence for *Trypanosoma brucei* parasites (61).

Beyond effects on microbial communities, glyphosate has broad physiological impacts

on insects, other arthropods, and vertebrates. While glyphosate was harmless to *Lepthyphantes*

tenuis spiders, it changed behavior and increased mortality in *Pardosa milvina* and *Neoscona*

theisi spiders (62–64). Glyphosate reduced learning in *Aedes aegypti* mosquitoes (65) and in

honeybees reduced survival and caused learning defects associated with feeding, homing, and

flight behaviors (66,67). Glyphosate and AMPA delayed development and reduced survival of

the arthropod *Daphnia magna* (68). Glyphosate induces oxidative stress and damage in many

organisms, including *D*. *magna*, insects (fruit flies), amphibians (African clawed frog, European

green toad, marsh frog), fish (brown trout, spotted snakehead fish), and mammals (rats) (69–

74), often linked with lipid peroxidation and expression of antioxidant defenses (catalase,

glutathione, and superoxide dismutase). In human erythrocytes, glyphosate and AMPA mixtures

increase ROS production (75,76).

Given the melanin-inhibitory properties of glyphosate in fungi (26), we examined the

roles of glyphosate and AMPA as inhibitors of insect melanization and phenoloxidase. We used

two distinct insect models that both rely on melanin-based immunity: *Galleria mellonella* – a

species of wax moths (Lepidoptera: Pyralidae), and *Anopheles gambiae* – a mosquito vector of

malaria (Diptera: Culicidae). Considering melanin's importance in insect immunity, we evaluated

glyphosate's effects on *G. mellonella* susceptibility to the pathogenic fungus *C. neoformans,* and

on *A. gambiae* survival and susceptibility to the malaria parasite *Plasmodium falciparum* as

measured by parasite oocyst burden. Additionally, we evaluated glyphosate's mechanism of

melanin inhibition using L-DOPA auto-oxidation and mushroom tyrosinase-mediated oxidation

models. Mushroom tyrosinase is commercially available and produces melanin in a similar

mechanism as insect phenoloxidase. The purified enzyme and L-DOPA auto-oxidation allowed

us to take a controlled step-by-step biochemical approach to show that glyphosate inhibits

melanization by disrupting oxidative balance.

RESULTS

Glyphosate and AMPA Inhibit *Galleria mellonella* Phenoloxidase Activity

In insects, phenoloxidases are activated by serine proteases upon wounding or infection,

thus triggering melanin production to either clot a wound or restrict a pathogen (77). To

investigate whether glyphosate inhibited insect melanogenesis, we used two models: *G.*

mellonella wax moth larvae, and *A. gambiae* adult mosquitoes, a main vector of malaria.

In an *ex vivo* analysis using *G. mellonella* hemolymph, we found that glyphosate inhibited

phenoloxidase activity in a dose dependent manner, without addition of exogenous substrate

(Fig. 1A). Similar results were found with addition of a broad-spectrum protease inhibitor, which

was used to control for continued activation of phenoloxidase, glyphosate-induced cellular

responses, and/or off-target effects on other components of the phenoloxidase cascade

(Supplementary Fig. 1A). We also saw similar inhibition of phenoloxidase activity with the

addition of exogenous L-DOPA; however, in these experiments there was only a modest

enhancement of phenoloxidase activity with lower glyphosate concentrations followed by

striking inhibition at higher concentrations (Supplementary Fig 1B). Importantly, glyphosate did

not impact hemocyte viability, as measured by trypan blue exclusion (Supplementary Fig. 1C).

Aminomethylphosphonic acid (AMPA), a primary breakdown product of glyphosate that

accumulates in the environment (45), inhibited melanization similarly to glyphosate using *G.*

mellonella hemolymph and a commercially available mushroom tyrosinase (Fig. 1B,

Supplementary Fig 1D). These data show that glyphosate inhibits melanization in insects similar

to what has been previously shown in fungi (26), thus indicating that glyphosate interferes with

major melanin-based processes in at least two kingdoms of life.

Glyphosate Alters *G. mellonella* Susceptibility to Infection

Next, we sought to determine whether glyphosate increased *in vivo* susceptibility of

Galleria mellonella larvae to foreign organisms. We injected *G. mellonella* final instar larvae with

2 µg of glyphosate (~8-12 ng/mg per larvae) followed by infection with *C. neoformans* or a mock

infection. The two mock infected groups, glyphosate-treated and phosphate-buffered saline

(PBS)-treated exhibited similar survival. However, in the infected groups, the glyphosate-treated

larvae died faster compared to the PBS-treated controls (Gehan-Breslow-Wilcoxen test, p =

0.013) (Fig. 1C). A similar, but non-significant, trend was seen with a *C. neoformans lac1Δ*

strain (Supplementary Fig. 1E). The *lac1Δ* strain is unable to produce melanin, an important

virulence factor in *C. neoformans* pathogenesis. This strain is less virulent in the *G. mellonella*

model (78), potentially contributing to the lack of significant differences between the glyphosate

and PBS-treated groups**.**

The decreased survival of the glyphosate-treated group infected with *C. neoformans*

was correlated with smaller melanized particles within nodules formed during infection (*in vivo*),

as compared to the PBS-treated group infected with *C. neoformans* (Fig 1D). For these

experiments, we infected and drugged *G. mellonella* as we do during normal infections, then

collected the hemolymph 24 hours later and imaged the nodules and aggregates that formed *in*

vivo. The PBS-treated non-infected group had smaller or virtually no melanized structures. In

two of three replicates, the PBS-treated infected group had more melanized structures than the

glyphosate-treated group (Supplementary Fig 1G-I). Further, in the glyphosate-treated infected

group we observed more *C. neoformans* cells, and those within nodules displayed lower

degrees of melanin encapsulation compared to PBS-treated larvae (Chi-squared test, p =

0.0034) (Fig 1E). The scoring was based on a system devised with 0 representing no melanin

encapsulation of the yeast cell to 4 being the most melanin encapsulation, as depicted in (Fig

1F). The one-time treatment with glyphosate did not disrupt time to larval pupation, a process

mediated by laccases and phenoloxidases (Supplementary Fig. 1F). These data suggest a

direct correlation between glyphosate treatment, and increased susceptibility of *G. mellonella* to

infection caused by decreased melanin-based immune response (nodulation).

Glyphosate Alters *A. gambiae* Phenoloxidases, Susceptibility to Infection with Malaria

Parasites, and Survival

To ascertain the impact of glyphosate on *Anopheles gambiae* mosquito melanization,

we measured the phenoloxidase activity in whole-body mosquito homogenate following the

addition of glyphosate. Similar to our results with *G. mellonella*, glyphosate inhibited

phenoloxidase activity of *A. gambiae* homogenate in a dose-dependent manner (Fig. 2A).

To investigate whether glyphosate rendered mosquitoes more susceptible to infection

with the human malaria parasite *P. falciparum*, adult female mosquitoes were fed on 10% sugar

solution supplemented with glyphosate at different concentrations for 5 days and then given a *P.*

falciparum-infected blood meal. Parasite burden was assessed through enumeration of the

Plasmodium oocyst stage at 8 days post infection (DPI). Glyphosate-fed mosquitoes had higher

oocyst burdens with an overall non-significant trend of increasing oocyst burden with increasing

dose of glyphosate (Fig. 2B). However, we observed a sharp decline in parasite burden in the

10 mM treated group, which was likely due to the increased mortality of this group (Fig. 2C)

resulting in few surviving mosquitoes to assess the intensity of infection. In a low *P. falciparum*

infection intensity assay (Supplementary Fig. 2A), we observed that glyphosate-treated groups

are more likely to be infected than control groups. This is important, as lower parasite burdens

are more reminiscent of infections in field conditions in malaria endemic regions (79–82).

Sugar preparations with glyphosate at environmentally relevant concentrations were

given to *A. gambiae* mosquitoes to ascertain the herbicide's effect on the mosquito's lifespan.

Compared to control mosquitoes (sugar-fed without glyphosate), mosquitoes given low

glyphosate doses (30 to 300 µM) showed statistically significant improved survival, while those

fed higher doses of glyphosate (1 to 10 mM) had equal or decreased survival, with the 10 mM

glyphosate-exposed group exhibiting significantly decreased survival (Fig. 2C). Additionally, we

used a Cox Mixed Effects Model to account for fixed and random effects, and calculated the

Hazard Ratios for each of the treatment groups (Fig. 2D). Hazard Ratios<1 indicate a reduced

risk of death compared to the control while Hazard Ratios>1 indicate enhanced risk of death

compared to the control group. With this model, the mosquitoes treated with lower glyphosate

concentrations had a Hazard Ratio less than 1, those treated with 1 and 3 mM glyphosate had

Hazard Ratios similar to 1, and the 10 mM-treated mosquitoes had a Hazard Ratio significantly

greater than 1. These results suggest that glyphosate could have bimodal effects on mosquito

health.

We also measured the impact of glyphosate on the mosquito cuticle and body size.

There was no discernable difference in *A. gambiae* cuticle pigmentation after 5 days of

treatment with 1 mM glyphosate in 10% sucrose from days 3 to 8 post-emergence, as

measured by mean gray value of the ventral abdomen (Supplementary Fig. 2C). This was

expected due to the typical expression of cuticular laccases being largely at the timing of

pupation and the first three days post-emergence (23,24). Additionally, there were no difference

in wing length, a proxy for adult size, between the glyphosate-treated and untreated adult

mosquitoes (Supplementary Fig. 2D). Altogether, this suggests that the observed increased

parasite burden in glyphosate-treated mosquitoes cannot be explained merely by broader

impacts of glyphosate on mosquito health.

Glyphosate Alters the Composition, but Not Density, of the *A. gambiae* Midgut Microbiota

A. gambiae midgut microbiota can influence *Plasmodium* infection by modulating the

mosquito's innate immune system and hence affecting parasite viability (83–86). We

investigated whether glyphosate had detrimental effects or influence on the *A. gambiae*

microbiota. Colony Forming Unit (CFU) counts from cultures of midgut homogenates grown on

LB agar demonstrated that glyphosate treatment did not affect total number of culturable gut

bacteria (Fig. 3A), though this method would miss any impacts on microbes that were not

readily cultured by these methods. To complement our culture-dependent analysis and provide

insight on microbiota community composition, we compared the total 16S rRNA composition of

the midgut microbiota with and without glyphosate treatment. Glyphosate treatment altered

microbiota composition, with a noted decrease in the relative abundance of Enterobacteriaceae

and an increase in relative *Asaia spp.* populations (Fig. 3B). We did not observe a dose-

dependent impact on composition, as the alpha diversity (a function of the number of bacterial

taxa) of mosquitos exposed to glyphosate was similar to controls (Fig. 3C). However,

community composition was perturbed by treatment with glyphosate, and glyphosate-treated

groups and controls form two separate clusters in principal coordinates analysis as measured

by Bray-Curtis dissimilarity (Fig. 3D). These differences suggest a shift in beta diversity

(prevalence of each bacterial taxon), and therefore a difference between the microbial

communities of mosquitoes exposed to glyphosate versus untreated controls.

Glyphosate Inhibits Production of Dopaquinone, Dopachrome, and Melanin

To understand how glyphosate inhibited melanization, we evaluated the formation of

melanin intermediates in a stepwise manner using a commercially available fungal tyrosinase

and the melanin precursor L-DOPA (2 mM). Although this tyrosinase differs from insect

phenoloxidase, the melanization reaction in these systems follows the same Mason-Raper

pathway (Figure 4A) (87,88) and thus can be used to explore the mechanism of glyphosate

inhibition. The first step of the reaction involves L-DOPA oxidation to dopaquinone (DQ)

enzymatically or spontaneously (89). We found that glyphosate inhibited the dopaquinone

production in a dose-dependent manner (Fig. 4B). This inhibition was observed for both

tyrosinase-mediated and auto-oxidation-mediated production of dopaquinone. The slopes of

inhibition in the auto-oxidation and tyrosinase-mediated oxidation were similar. This indicated

that the tyrosinase reaction dopaquinone levels would remain unchanged by glyphosate

treatment if the inhibition of "background" auto-oxidation dopaquinone production were taken

into consideration. These results suggested that dopaquinone inhibition was primarily rooted in

preventing the oxidation of L-DOPA independent of tyrosinase.

Dopaquinone spontaneously cyclizes to form cyclodopa, which then undergoes a redox

exchange with another dopaquinone molecule to form one molecule of dopachrome and one

reformed molecule of L-DOPA. Dopachrome is a pink-orange melanin intermediate that has an

absorbance maximum at 475 nm. Dopachrome is a useful proxy product for tyrosinase-

mediated reaction kinetics and evaluating the melanization reactions and redox exchange (90).

The rate of dopachrome formation and the amount of dopachrome produced were determined

by measuring changes in absorbance during a reaction between L-DOPA and tyrosinase. There

was a strong dose-dependent inhibition of dopachrome formation with glyphosate (Fig. 4C),

implying that the compound's inhibitory effects were upstream of dopachrome.

We tracked the reaction over 5 d to confirm inhibition of melanin synthesis. Glyphosate

inhibited the production of a black pigment dose-dependently, as measured by the absorbance

of the tyrosinase reaction on Day 5 (Fig. 4D). Interestingly, glyphosate also inhibited the

formation of pigment that derives from auto-oxidation of L-DOPA (Fig. 4D). This implies that

glyphosate inhibited pigment production non-enzymatically.

Phosphate-Containing Compounds Inhibited Melanization Similarly to Glyphosate

To gain insight into the chemical features of glyphosate that inhibited melanogenesis we

assayed several structurally similar compounds using the same *in vitro* mushroom tyrosinase

assay. To test the effect of the amino acid functional group, we compared glyphosate alongside

its non-phosphate analog, glycine. We also tested the inhibitory effects of phosphoserine and

serine on melanin production. Phosphoserine inhibited dopaquinone, dopachrome, and melanin

formation to nearly the same extent as glyphosate (Fig. 5A-C). In contrast, neither glycine nor

serine inhibited dopaquinone, dopachrome, or overall melanin formation (Fig. 5A-C). We tested

the inhibitory effects of other phosphate-containing compounds including organophosphates

(phosphonoacetic acid), phosphoesters (pyrophosphate), and phosphoric acid. All of the

phosphate-containing compounds inhibited dopaquinone production (Fig. 5A) and dopachrome

formation (Fig. 5B) in a manner nearly identical to glyphosate, but differed slightly from each

other in melanin inhibition (Fig 5C).

Similar to glyphosate, these compounds all inhibited auto-oxidation of L-DOPA

comparably to their inhibition of tyrosinase-mediated melanin production (Fig. 5E). This further

illustrates that glyphosate and similar phosphate-containing compounds inhibit melanin in a non-

enzymatic fashion. These data suggest that the phosphate functional groups of these

compounds may be responsible for the melanin-inhibitory properties.

Glyphosate Does Not React With L-DOPA or Inhibit Tyrosinase Directly

We considered the possibility that glyphosate inhibited melanogenesis and dopaquinone

production by reacting with the L-DOPA substrate. To measure the reaction between these

compounds, we analyzed mixtures of L-DOPA and glyphosate by ^{1}H-NMR and ^{31}P-NMR. We

found no evidence of interaction between the two compounds based on peak shifts of hydrogen

and phosphorous at both high (60 mM glyphosate and 20 mM L-DOPA) and low concentrations

(6 mM glyphosate and 5 mM L-DOPA) (Supplementary Fig. 4).

If glyphosate was inhibiting melanin production through the formation of a covalent bond

with tyrosinase, the inhibition should be irreversible. To test this, we treated 20 µg/ml tyrosinase

with 5.63 mg/ml (33.33 mM) glyphosate and removed the glyphosate by dialysis. The

glyphosate-treated enzyme had similar activity to the control (Fig. 6A), making a strong case

against a mechanism whereby glyphosate inhibited melanogenesis through irreversible

inhibition of tyrosinase. Instead, analysis of the tyrosinase reaction by Michaelis-Menten kinetics

assay with L-DOPA and glyphosate suggested that glyphosate is a non-competitive inhibitor of

melanin and dopachrome production (Fig. 6B). Further, we tested tyrosinase activity as a

function of enzyme concentration with and without glyphosate and constant concentration of L-

DOPA. We found that the slope of the glyphosate-treated enzyme is lower than the water-

treated control (Fig. 6C). This indicates that glyphosate-mediated inhibition is reversible (91,92).

Given that our findings showed that glyphosate inhibits auto-oxidation and tyrosinase-mediated

oxidation, we believe that the reversible inhibition is due to glyphosate interfering with the L-

DOPA substrate's ability *to be* oxidized rather than the enzyme's ability *to* oxidize. This could be

represented by the following where *E* represents tyrosinase, *S* represents L-DOPA, *I* represents

glyphosate, and *P* represents dopaquinone/melanin:

315   Normal Enzymatic Reaction: $E + S \rightleftharpoons ES \longrightarrow P$

316   Inhibited Enzymatic Reaction: $E + S + I \rightleftharpoons E + SI \dashv P$

Copper ions are important for tyrosinase activity. Since glyphosate is a metal chelator[,]

(93,94), we evaluated whether glyphosate's inhibitory effect was due to this property. We added

copper ions to the L-DOPA and tyrosinase reaction to rescue the glyphosate inhibition. We

performed the experiment with eight concentrations of copper (II) sulfate for each of the eight

glyphosate concentrations. In general, the addition of copper did not rescue the glyphosate

dependent inhibition of melanin (Fig. 6D). However, low concentrations of copper (6.25 - 25 µM)

increased tyrosinase activity, while high concentrations of copper (50 - 400 μM) reduced activity,

indicating low copper can boost enzyme activity, while higher concentrations inhibit the reaction.

However, this hormesis-like effect was not observed at increasing glyphosate concentrations

(Supplementary Fig. 5). This result indicates that glyphosate's ability to chelate copper ions

could have a protective effect in high copper environments, which would otherwise lead to

negative effects on enzymatic activity and other biological processes. Similar results have been

previously seen in *Eisenia fetida* earthworms exposed to high copper conditions in soil (95).

Glyphosate contamination of copper-rich soil reduced the detrimental effects of the metal's

toxicity, presumably due to the glyphosate's copper chelation properties (95).

Glyphosate Affects the Oxidative Properties of Melanogenesis

Melanogenesis is dependent on the spontaneous radicalization of quinone intermediates

(96). Dopaquinone radicals and cyclodopa undergo a radical-mediated redox exchange that

converts cyclodopa into dopachrome and dopaquinone into L-DOPA. Further downstream, ROS

catalyze the polymerization of dihydroxyindole into eumelanin. Glyphosate's inhibitory effect

could be due to a role as a free-radical scavenger or antioxidant. Since the inhibitory

compounds blocked spontaneous oxidation of L-DOPA (Fig. 5E), they are antioxidants. To

measure this radical-quenching ability we used an ABTS assay in which ABTS radicals are

blue, but when quenched the solution becomes colorless. The degree of discoloration is a proxy

for radical concentration and antioxidant strength. Glyphosate quenched the ABTS radical to

some degree, but only after several hours of reaction (Supplementary Fig. 6A), which did not

occur with the other inhibitory phosphate-group containing compounds evaluated (Fig. 7A). This

indicates that direct free-radical scavenging may not be the primary mechanism of melanin

inhibition for glyphosate.

Phosphoric acid is a well-known synergist that boosts the antioxidant properties of

phenolic compounds. Phosphoric acid, and other synergists such as citric acid, malic acid, and

tartaric acid do not directly quench free radicals themselves, but instead work by regenerating

antioxidants, thus becoming "sacrificially oxidized", or chelating metal ions in solution (97,98).

Alternatively, glyphosate could be reacting with existing antioxidants to strengthen and/or

regenerate them into "active" form. In this instance, the glyphosate would be bolstering the

antioxidant properties of L-DOPA.

We observed that the synergist citric acid inhibited melanization similarly to glyphosate

and phosphoric acid (Fig. 7B,C). The addition of glyphosate, phosphoserine, and phosphoric

acid enhanced the antioxidant properties of L-DOPA in an ABTS assay in a similar manner as

citric acid (Fig. 7D). This suggests that glyphosate may act as an inhibitor via antioxidant

synergism. The synergy is the ratio of the quenching capacity of the L-DOPA and the

compounds alone to the quenching capacity of L-DOPA combined with the compound. The

lower this ratio, the more synergistic the compounds are with L-DOPA (Supplementary Fig. 6B).

These values indicate that the inhibitory compounds are synergistic, whereas the non-inhibitory

glycine and serine are not as synergistic.

The inhibition of melanin was independent of the L-DOPA to glyphosate ratio, and

glyphosate's IC_{50} is ~1 mM regardless of L-DOPA concentration (Supplementary Fig. 7). This

could be explained by a general antioxidant effect on solution.

Glyphosate Alters the Oxidation-Reduction Potential of the System

L-DOPA is a more effective antioxidant when it is oxidized or radicalized, and has a

better ability to form adducts with other radicals (99). Since glyphosate is acting as a synergistic

antioxidant, it may be driving L-DOPA oxidation and possibly radicalization in which L-DOPA

scavenges radicals better. This has the potential to disrupt melanin synthesis by stopping the

spontaneity of redox exchange and dopaquinone formation.

To investigate whether the addition of glyphosate changed the oxidation properties of L-

DOPA, we used cyclic voltammetry – a technique to measure the electrochemical properties of

solutions and previously used to study quinone electrochemistry (100,101). Voltammetry

performed on L-DOPA solutions with glyphosate showed dose-dependent shifts towards a

negative potential (Fig. 7E,H) in peaks that corresponded to L-DOPA oxidation (102) (**Peak 1**).

We validated these as L-DOPA oxidation peaks by performing voltammetry on various L-DOPA

concentrations **(**Supplementary Fig. 8A). The peak shift towards negative potentials indicates

the L-DOPA was oxidized more easily and had less ability to be an oxidant, similar to the

negative potential shifts associated with alkaline pH and increased oxidation (103). We

controlled for any pH-dependent peak shifts by adjusting each solution to pH 6.00 prior to

measurement. Decreased oxidizing power can lead to significant effects, as melanin

biosynthesis is reliant upon catechol oxidation and high redox potentials of quinones.

Interestingly, with increased glyphosate, the L-DOPA solution had a lower current intensity

associated with the reduction of dopaquinone to L-DOPA (**Peak 2**). In cyclic voltammetry,

smaller peaks indicate that less of the compound is oxidized or reduced. The decreased **Peak 2**

current became virtually non-existent with increasing glyphosate concentrations (Fig. 7E,H,I).

This implies that dopaquinone, represented by **Peak 2**, is either not being formed during L-

DOPA oxidation or cannot be reduced back into L-DOPA. These data indicate that the redox

cycling steps of melanization are halted due to the inability of dopaquinone to be reduced into L-

DOPA. This could also indicate that while L-DOPA was oxidized more in the presence of

glyphosate, it may form a non-dopaquinone product - either a radical-mediated dimer with itself

or a semiquinone.

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DISCUSSION

We investigated the effect of glyphosate on melanin production in two species of insects,

Galleria mellonella and *Anopheles gambiae*, and found that both glyphosate and its major

metabolite AMPA were inhibitors of insect phenoloxidase and melanization. Although

glyphosate and AMPA are relatively weak inhibitors of these insects' melanization, the inhibitory

concentrations are relevant in the environment given the vast amounts used in agriculture, their

environmental stabilities, and the high potential for insect-herbicide interactions. Therefore,

glyphosate has a high potential to influence key insect physiological systems. We observed that

glyphosate enhanced the susceptibility to infection of two phylogenetically distinct insects, *G.*

mellonella and *A. gambiae*. This raises concerns and the suggestion that glyphosate may

interfere broadly with insect immunity through its effects on melanin-based defenses. Analysis

of *in vitro* tyrosinase and auto-oxidation models revealed that glyphosate inhibited melanization

by acting as a synergistic antioxidant and disrupting redox cycling. Overall, our findings provide

new insights on the complex reaction and suggest potential harmful effects of this herbicide on

non-target organisms, including some insects that may be important to ecosystem stability, and

already in peril due to the threat of an "insect apocalypse".

G. mellonella treated with glyphosate were more susceptible to infection with *C.*

neoformans. Glyphosate treatment was associated with reduced size of melanized nodules in

the hemolymph following infection with *C. neoformans*. Two of three replicates showed

significantly reduced numbers of melanized nodules in the glyphosate-treated infections.

Nodules are primarily composed of hemocyte aggregates, released immune factors, and

melanin encapsulation of the pathogen, which function together to kill invading pathogens (22).

Altogether, these data suggest that glyphosate weakened the melanin-based immune response

of *G. mellonella,* which could have grave implications for host defense. *Galleria* are members of

the order Lepidoptera (moths and butterflies), which represent up 10% of known species on

Earth. Interactions with glyphosate in the soil, on plants during pollination, or ingested through

herbivory could contribute to immunocompromised lepidopteran populations. Glyphosate's

effects on immunity in insects could compound a controversial and pre-existing problem of

declines in Lepidopteran biomass in recent decades (25,104–107).

Like our observations with *G. mellonella*, glyphosate made the *A. gambiae* mosquito

more susceptible to *P. falciparum* parasite infection, the primary agent of human malaria in

Africa. However, melanization is not considered the primary anti-*P. falciparum* immune

response in this malaria model (108). The increased susceptibility of *A. gambiae* to *P.*

falciparum could be due to broader alterations of mosquito immune defenses, or disruption of

non-melanin roles of catecholamine oxidation and phenoloxidase in insect immunity including

the production of ROS, cytotoxic intermediates, and pathogen lysis (3,13,109). Importantly, we

observed that even when infections of *A. gambiae* with *P. falciparum* resulted in an overall low

to no parasite burden in control-treated groups, glyphosate-treated groups exhibited a higher

infection burden and prevalence. This is notable because *Plasmodium* oocyte development

within the mosquito is a major bottleneck to successful vector competence in nature (110). If a

mosquito can prevent oocyst formation, there is no transmission of malaria to humans. The

numbers of oocysts from these low parasite burden experiments are in line with the normal

burden's found in natural field infection models (111,112). Our data may indicate that

mosquitoes exposed to glyphosate were less able to control *Plasmodium* infection they would

have otherwise resisted, thereby becoming potentially better vectors for malaria. Overall our

results raise concerns for public health and malaria control initiatives in regions in which malaria

is endemic and where there is increasing use of glyphosate, including areas of Latin America,

Sub-Saharan Africa, and Asia.

Our data revealed that uninfected adult female mosquitoes treated with glyphosate

displayed a hormesis-like dose-dependent effect when measuring survival outcomes. Survival

increased at low doses of glyphosate compared to the control. This greater longevity may be

due in part to reduced basal damage from host defense mechanisms that normally occur during

melanin formation, and/or altered gut microbiota. In contrast, mosquitoes exposed to high

concentrations of glyphosate showed decreased survival. These data suggest the broader

notion that glyphosate could have varied and complex outcomes on vector competence

depending on its concentration in the environment. The low-concentration glyphosate

treatments resulted in longer-lived, yet immunosuppressed, mosquitoes that were slightly more

susceptible to infection with *P. falciparum,* whereas short-lived high glyphosate treated

mosquitoes were much more susceptible to *P. falciparum.* Further, while the 10 mM-treated

mosquitoes had the worst survival outcome, the mosquitoes that survived the drugging showed

low susceptibility to *P. falciparum* infection. These observations suggest a potentially interesting

effect whereby very high concentrations of glyphosate reduce mosquito survival, but bolster the

immune system or general physiology of survivors, which then allows them to resist *P.*

falciparum infection with greater success. Alternatively, very high glyphosate treatment could be

selecting for mosquitoes within the population more resistant to *P. falciparum* infection.

Our analyses of *A. gambiae* midgut microbiota indicated that glyphosate did not impact

A. gambiae midgut culturable bacterial density; although the herbicide did perturb midgut

microbiota composition in a non-dose dependent manner. More specifically, glyphosate altered

diversity of the microbial community, and glyphosate-treated mosquitoes exhibited diminished

Enterobacteriaceae and expanded *Asaia spp.* populations. The presence of some

Enterobacteriacae, including the common insectary contaminant *Serratia marcescens,* in

Anopheles spp. midguts is associated with lower susceptibility to *Plasmodium spp.* infection

(113,114). This effect is observed quantitatively by the significantly different prevalence of

individual bacterial taxons (beta diversity) between the glyphosate and control-treated

microbiota, while there is an overall unchanged number of bacterial taxa present (alpha

diversity). Beta diversity analysis indicates that microbial communities associated with

glyphosate-treated mosquitoes cluster together and are different than those from control

mosquito communities.

Our results are consistent with reports that glyphosate perturbs the microbiota of

honeybees that makes them more susceptible to infection (59). Our data suggests that while

glyphosate may perturb the microbiota and affect immunity as previously described (59,60), it

can also inhibit melanization which is a critical part of insect immune defense. We do not see a

dose-dependent effect of glyphosate on the microbiota composition, but we do see a dose-

dependent effect on mosquito susceptibility to *Plasmodium* infection; this indicates that the

enhanced susceptibility might be unrelated to microbiome perturbations. These mechanisms of

susceptibility are not mutually exclusive and could be additive to weaken insect health.

Additionally, while AMPA does not to disrupt the microbiota of honeybees (60), we show it can

inhibit melanization of *G. mellonella* phenoloxidase and mushroom tyrosinase. A recent study in

Apis cerana cerana honeybees indicate that glyphosate-based herbicide treatment increases

expression of wound and defense genes, including those related to melanization (115).

Interestingly, this study also showed that glyphosate feeding decreased the expression of many

odorant binding proteins, which have been shown to mediate the melanization response in both

Tsetse fly (*Glossina morsitans morsitans*) and *Drosophila melanogaster* (116) This suggests

complex regulation of melanization following treatment with glyphosate-based herbicide,

including the possibility of increased melanin-related gene expression as a compensation for

glyphosate's inhibitory effects. Additionally, other surfactants and components of the

commercial herbicide formulation used could trigger damage and immune gene expression.

Melanins and phenoloxidases are involved in other physiological functions in insects

including proper pupation, and cuticle and eggshell development. In our experiments, which

involved single dosing or short duration of feeding, we did not detect a difference in coloration of

adult mosquito treated with glyphosate, nor a defect in *G. mellonella* pupation following

glyphosate treatments of final instar larvae. However, we cannot rule out that longer glyphosate

exposure or feeding throughout the lifecycle would effect these functions. If such effects

happen, they will only compound the effects of glyphosate on melanin-based immunity and

insect physiology.

We sought to understand the mechanism of melanization inhibition by glyphosate. The

process of melanization is highly dependent upon oxidation and redox cycling between

catechols and quinones. The melanin production process is halted if the oxidizing ability and the

redox potentials are altered. Melanization begins with the conversion of L-DOPA into

dopaquinone through enzymatic or spontaneous oxidation of L-DOPA, followed by redox cycling

that results in dopachrome formation, and subsequently melanin polymerization. Glyphosate

inhibited formation of dopaquinone and melanin pigment mediated by both tyrosinase and auto-

oxidation, which strongly suggests that glyphosate inhibits L-DOPA oxidation in an enzyme

independent manner. We found that other phosphate-containing compounds inhibited

melanization in a similar manner including phosphoserine, phosphoacetic acid, pyrophosphate

and phosphoric acid. This is in line with literature reports that other aminophosphonic acids

inhibit fungal eumelanin in the human pathogen *Aspergilius flavus* (117). Incidentally, this class

of compounds is patented for use in human cosmetics, and are marketed as solutions to inhibit

melanogenesis in the skin (118,119).

We found no evidence that glyphosate irreversibly inhibited tyrosinase activity or directly

interfered with enzyme function. Addition of copper ions did not rescue the inhibition, indicating

that the copper-based catalytic core of tyrosinase in not disrupted by glyphosate. Interestingly,

low copper increased tyrosinase activity and high doses reduced activity. However, copper had

minimal effects on tyrosinase activity during high glyphosate concentrations. It appears that

glyphosate, possibly through chelation, acts as a "buffer" of copper ions and can reduce the

metal's harmful effects, similar to previous findings concerning the toxicity of high-copper soil to

earthworms following glyphosate treatment (95). This could have broader implications for

melanogenesis in nature, where some fungi use copper as a signal to upregulate melanin-

producing enzymes (120), and thus copper ion sequestration could reduce melanin production.

We examined the ability of glyphosate and the other compounds to quench free radicals,

which are necessary to the melanization process. Of the inhibitors tested, only glyphosate had

radical-quenching activity, but this occurred relatively slowly compared to the typical timeframe

of antioxidant reactions reported in literature (121). This property is likely not the mechanism of

inhibition as phosphoserine has a similar structure and near identical inhibition of melanization

as glyphosate, yet no radical-quenching properties. While not a free-radical quencher,

phosphoric acid is a known antioxidant synergist – a class of compounds that enhance

antioxidant properties of phenolic compounds by chelating metals or reverting antioxidants into

their active states (98). Synergists like phosphoric acid, citric acid, malic acid, and alpha-

hydroxy acids are added to foods, medicines, and cosmetics at concentrations up to 10% as a

preservative due to their synergist effects on antioxidants (122). Glyphosate behaved similarly

to phosphoric acid and citric acid; citric acid inhibited melanization similarly to glyphosate and

phosphoric acid, suggesting an inhibition mechanism via antioxidant synergy. Additionally, we

report that glyphosate and other inhibitors have synergistic effects on the antioxidant properties

of L-DOPA. L-DOPA's antioxidant properties derive from its reduction back to a normal state

from an oxidized state, or a radical-mediated adduction reaction with the oxidized compounds in

solution. Since glyphosate makes L-DOPA a more efficient antioxidant, glyphosate thus alters

the oxidative balance of L-DOPA and/or produces a buildup of radical or semiquinone

intermediates.

Consistent with these findings of antioxidant synergy, cyclic voltammetry revealed that

glyphosate decreased the L-DOPA-Dopaquinone redox potential. Hence, L-DOPA becomes

both a weaker oxidizing agent and a stronger reducing agent (antioxidant) and is more prone to

oxidation in the presence of the herbicide. Glyphosate decreased dopaquinone reduction in a

dose dependent fashion indicating that dopaquinone cannot be reduced or is not produced

following L-DOPA oxidation. A lack of dopaquinone could indicate that glyphosate causes

oxidized L-DOPA semiquinone intermediates to remain stable or react with each other and form

L-DOPA dimers. On the other hand, if dopaquinone cannot be reduced into L-DOPA,

melaninization becomes unfavorable as redox exchange could not occur. These changes in

voltammogram do not appear when the L-DOPA solution is treated with 16 mM glycine, but did

occur with citric acid. This further supports that glyphosate is acting as a synergistic antioxidant

and prevents the redox-dependent melaninization.

Our findings investigating glyphosate's mechanism of melanin inhibition points to

disruption of oxidative balance and redox cycling which may result in the buildup of toxic

oxidative intermediates. Previous studies evaluating glyphosate's impact on organisms show

that the herbicide increases oxidative stress, lipid peroxidation, and antioxidant responses in

bacteria, plants, arthropods, fish, amphibians, rats, and human red blood cells (69–76). These

data bolster our findings that glyphosate promotes oxidation in phenolic compounds like L-

DOPA and inhibits clearance of oxidative stress. Understanding the mechanisms by which

compounds such as glyphosate might impact insect biomass and contribute to a potential insect

decline is important, as they have both direct and indirect impacts on human health.

Glyphosate's interference with melanization could have considerable environmental

impact given its stability and wide concentration range, from over 50 mM at time and at site of

application to under 1 nM in runoffs from application sites (33,39,123). At higher

concentrations, glyphosate could inhibit melanin production in some insects, thus rendering

them more susceptible to pathogens due to reduced immune competence. This suggests

protean consequences for human health ranging from ecosystem disruption to altered vector

competency of lethal human pathogens and increased malaria transmission in endemic regions

that use glyphosate-based herbicides in agriculture. Importantly, we provide evidence that

glyphosate enhances *A. gambiae* susceptibility to the human malaria parasite, which could

potentially make it a better vector for transmitting disease to humans. Our data in *Galleria* and

Anopheles can perhaps be extrapolated to other lepidopteran (moth and butterfly) and dipteran

(fly) species with additional importance to the environment.

In summary, our results suggest that glyphosate interferes with melanization in two

insect species, through a mechanism involving altering the redox potential of melanin

polymerization reaction. This phenomenon is concerning because of the importance of

melanization in insect immunity. A strong immune response is vital for insect survival, and

disruption of their immune function, including the inhibition of melanization, could be disastrous

for these animals. Insects are pivotal members of the world's ecosystems, essential to

maintaining proper function, and they ensure human food security. Yet, certain data indicates a

drop in insect biomass over recent decades, a phenomenon that has been called the "insect

apocalypse" (25,104,105,124,125). Although this view has been questioned regarding the true

extent and possible causes of the insect population declines (126–130), our results suggest that

glyphosate use as a mechanism by which insect immunity can be undermined by human

activities.

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AUTHOR CONTRIBUTIONS

D.F.Q.S. contributed to conceptualization, methodology, formal analysis, investigation, data

curation, writing of the original draft and subsequent editing, data visualization, and project

administration. E.C. contributed to conceptualization, methodology, investigation, resources,

reviewing and editing of manuscript, supervision, project administration, and funding acquisition.

R.T. contributed to conceptualization, methodology, resources, reviewing and editing of

manuscript, and supervision. A.J.B. contributed to conceptualization, methodology, software

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visualization, and project administration. Y.D. contributed to methodology and reviewing and

editing of manuscript. G.D contributed to reviewing and editing of manuscript, resources, and

funding acquisition. N.A.B. contributed to conceptualization, methodology, investigation,

resources, reviewing and editing of manuscript, supervision, project administration, and funding

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administration, and funding acquisition.

DECLARATION OF COMPETING INTERESTS

The authors declare no competing interests.

METHODS

Biological materials

Galleria mellonella larvae were obtained through Vanderhorst Wholesale Inc, St. Marys,

Ohio, USA. *Cryptococcus neoformans* strain H99 (serotype A) was kept frozen in 20% glycerol

stocks and subcultured into Sabouraud dextrose broth for 48 h at 30°C prior to each

experiment. The yeast cells were washed twice with PBS, counted using a hemocytometer

(Corning, Inc), and adjusted to 10^6 cells/ml.

Anopheles gambiae (Keele strain) mosquitoes were maintained on sugar solution at

27°C and 70% humidity with a 12 h light to dark cycle according to standard rearing condition.

Plasmodium falciparum NF54 (Walter Reed National Military Medical Center, Bethesda)

infectious gametocyte cultures were provided by the Johns Hopkins Malaria Research Institute

Parasite Core Facility and were diluted to 0.05% gametocytemia with naïve human blood before

feeding to the mosquitoes using an artificial glass membrane feeder as established in (Dong et

al., 2009) (83).

Compound and Dilution Preparation

Each compound, including the glyphosate (Millipore Sigma, Product #45521), was

prepared in 300 mM stock solution in Milli-water Q and brought to a pH of 5.5, and 20 µl of each

compound was serially diluted 1:2 in PBS (pH 7.4), with a compound-free control. When all

reaction components are added, the final concentrations of the compounds in these dilutions

were 33.33, 16.67, 8.33, 4.17, 2.08, 1.04, 0.52, and 0 mM.

Galleria mellonella **Hemolymph Extraction and Phenol Oxidase activity**

Healthy (active and cream-colored) larvae were cold anesthetized, punctured in their

proleg with 18G needle and pressure was applied to the larvae to promote bleeding of

hemolymph. Hemolymph was collected from larvae directly into an eppendorf tube.

Anticoagulants were not used as they might interfere with the melanization process.

For automelanization experiments, hemolymph was diluted 1:10 in PBS and mixed with

a pipette. Then, 160 µl of 1:10 hemolymph is added to 20 µl of glyphosate serially diluted in

PBS. The change in absorbance at 490 nm was recorded and analyzed as described above.

For experiments with L-DOPA, hemolymph was diluted 1:5 in PBS and mixed by pipette.

Experiments were performed as per the phenoloxidase activity assay in (Cornet, Gandon, and

Rivera, 2013 (131)) .

To test the effect of glyphosate on hemocytes viability, hemolymph was diluted 1:2 with

anticoagulation buffer (132), as melanization was not of importance for this experiment.

Hemocytes were pelleted and suspended in anticoagulation buffer. Glyphosate was added to an

aliquot of hemocytes in solution and incubated with mixing on a rocker at 30°C for 15 min.

Hemocyte viability was assessed by 0.02% trypan blue staining and enumeration of stained

(dead) versus unstained (alive) hemocytes with a hemocytometer.

Galleria mellonella **Infection and Survival**

Healthy final instar *G. mellonella* larvae weighing between 175 and 225 mg were

selected, and left starving overnight. Groups of larvae were injected with 10 µl of PBS or 10 µl of

1 mM sterile glyphosate in PBS. Larvae were monitored and left to recover for 5 h. Larvae were

then injected with 10 µl of sterile PBS or injected with 10^4 *Cryptococcus neoformans* yeast cells

per larva. Due to the low concentration of glyphosate administered to the larvae, their volume of

hemolymph, and their body volume, we believe the approximate concentration of glyphosate is

below the concentrations required to inhibit *C. neoformans* growth (26). *G. mellonella* larvae and

pupae were kept at 30°C and monitored daily for survival for 14 d. Survival was assessed by

movement upon stimulus with a pipette. See Supplementary Fig. 9a.

Melanization and Nodule Measurements

G. mellonella larvae were drugged and infected as described above in groups of 3 larvae

per condition. After 24 h, larval hemolymph was removed directly in anticoagulation buffer,

centrifuged at 10,000 *xg* for 5 min, and resuspended in coagulation buffer.

Brightfield microscopy images were randomly taken at 4x magnification, with 15-20

images taken per condition per replicate. These images were analyzed using Fiji (133) Particle

Analyzer function using with a threshold set between 0 and 120 mean gray value. Particle area

and numbers were calculated. Additional images were taken of nodules at 20x and 100x

magnification, the latter of which were used to manually score the degree of melanization of

fungal cells within the nodules. Statistical significance of differences between melanized particle

area was analyzed using a nested non-parametric Mann-Whitney-Wilcoxon rank test using the

nestedRanksTest package (*Version 0.2*, D.G. Scofield , 2014)(134) in R for R 4.0.2 GUI 1.72 for

Mac OS at https://www.r-project.org/ (R Core Team, 2020).

***Anopheles gambiae* Phenol oxidase activity**

Phenoloxidase activity assays were performed as previously described (135).

Experiments were done in biological triplicate with different batches of mosquitoes, as well as in

technical triplicate per biological replicate of 3 batches of 10 mosquitoes.

***Anopheles gambiae* Survival**

Adult female mosquitoes of *A. gambiae* Keele strain were raised on 10% sucrose for

three days post-emergence. On the third day, adult females were sorted into seven groups of 40

and placed into mesh-covered cardboard cups and provided a cotton ball with 10% sucrose

mixture with either 0 µM (Control), 30 µM, 100 µM, 300 µM, 1 mM, 3 mM, or 10 mM glyphosate.

The cotton balls were replaced every third day with new cotton balls and fresh

sucrose/glyphosate solutions. Mosquito death was monitored daily for 14 days. Experiments

were performed in three independent replicates, for a total of 120 mosquitoes in each treatment

group.

***Anopheles gambiae* Cuticle Pigmentation and Wing Size**

Adult female mosquitoes were drugged for 5 days as previously described. Mosquitoes

were cold euthanized and mounted dorsally on a slide with double-sided tape. Images of the

mosquito ventral abdomen were taken under a dissection microscope with constant exposure

and lighting conditions. Pigmentation was measured using Fiji software (133). The entire

abdomen of each mosquito was selected using a freehand selection tool, and the 8-bit mean

gray value was measured using the Measure tool. A measurement of 0 corresponds to a pure

black gray value, whereas 255 corresponds to a pure white gray value.

Following abdomen pigmentation measurements, mosquito bodies were removed, with

careful attention to keeping the wings remaining intact on the tape. Intact wings were imaged on

a microscope, and the length of the individual wing lengths were measured from tip-to-tip using

Fiji Measure tool.

Anopheles gambiae* infection with *Plasmodium falciparum

Adult female mosquitoes (3-4 d old) of *A. gambiae* Keele strain were sorted and drugged

as described above. On the fifth day of glyphosate exposure, mosquitoes were provided a blood

meal containing *P. falciparum.* Blood-fed engorged mosquitoes were sorted on ice and fed 10%

sucrose *ad libitum* for 8 d. Midguts were dissected and stained with 0.2% Mercurochrome

solution and oocysts were enumerated using a 20X objective with light microscopy. See

Supplementary Fig. 9b.

Anopheles gambiae **Midgut Microbiome Analysis**

Adult female mosquitoes (3-4 d old) of *A. gambiae* Keele strain were sorted and drugged

as described above. On the fifth day of glyphosate exposure, mosquitoes were sterilized in

ethanol for 2 minutes, washed, and dissected in sterile PBS. The midguts were removed, placed

in 500 µl sterile PBS on ice, homogenized, diluted, and plated on LB agar plates. Plates were

incubated at 30°C for three days and individual colonies were counted. Each experiment used

10-20 mosquitoes per condition, and the experiment was performed three independent times.

For the 16S rRNA sequencing studies, mosquitoes were reared, drugged, and then

midguts were dissected as described above, with five individual midguts per condition. DNA was

extracted from frozen mosquito samples using the Lucigen EpiCentre MasterPure DNA

extraction kit. The bacterial 16S rRNA gene was amplified by PCR, and sample-specific Illumina

adapters were ligated to the PCR products. PCR products from multiple samples were pooled

and sequenced on the Illumina MiSeq platform by the University of Connecticut MARS Facility.

Data were then analyzed using mothur (136) to construct contigs to align forward and reverse

reads, remove ambiguous bases and chimeric regions, align sequences to the Silva 16S V4

reference database, and cluster reads into 3% operational taxonomic units (OTUs). Sequences

derived from known contaminants were selectively removed. Alpha and beta diversity

measurements were performed using the Shannon diversity index and Bray-Curtis dissimilarity

distance respectively. Bray-Curtis distances were graphed on principal coordinates analysis

(PCoA) plots in two dimensions. Taxa and PCoA graphs were produced using

MicrobiomeAnalyst (137,138). See Supplementary Fig. 9c.

Dopaquinone Formation MBTH Assay

Quinones like dopaquinone are unstable and difficult to study directly; thus,

dopaquinone quantification relies on the formation of a stable adduct with MBTH (3-methyl-2-

benxothiaxolinone hydrazine) that forms a pigment that absorbs at 505 nm(139). This

absorption overlaps with the absorption of another melanin intermediate, dopachrome (Q), but is

not expected to interfere since dopaquinone reaction with MBTH prevents dopachrome

formation. Further, the molar absorbance coefficient for MBTH-Dopaquinone is more than 10

times higher (39,000 L/[mol cm]) than that of dopachrome (3,700 L/[mol cm]), and interference

from dopachrome would be relatively small.

MBTH reaction mixtures were prepared as previously described (139). This mixture is

warmed at 42°C to help solubilize the components. Then, 5 µl of 2 µg/ml Mushroom Tyrosinase

(Sigma, Product #T382) and 20 µl of 20 mM L-DOPA are added to the MBTH solution, and 160

µl of the solution is immediately added to each well containing compounds. The plate was read

at an absorbance of 505 nm for 30 min at 30°C, and read again at 1 h and overnight. The

dopaquinone levels are determined by the formation of the bright pink adduct between the

quinone and the MBTH.

Dopachrome and Melanin Measurements

Tyrosinase activity was determined as previously described (135), substituting

mushroom tyrosinase for phenoloxidase. The formation rate of dopachrome is measured as the

maximum velocity of this reaction, and the dopachrome levels are measured as the absorbance

at 490 nm after 30 min as the absorbance values plateau. Melanin levels are measured as the

absorbance at 490 nm after the reaction has continued for 5 d in the dark at room temperature.

Free-Radical Scavenging ABTS Assay

ABTS solution was prepared as previously described (140). To test the radical-

scavenging capability of the compounds, 10 µl of the compounds were serially diluted in a 96

well plate as previously described, and 90 µl of diluted ABTS was added to each well. The 734

nm absorbance was measured immediately, after 10 min, 1 and 2 h. In kinetics experiments,

absorbance readings were taken every two minutes for 5 h.

To measure the radical scavenging capacity of the synergistic compounds and L-DOPA

mixtures, ABTS was prepared and diluted in Milli-Q water. In each well, 5 μl of compound stocks

were added with either 5 μl of water or 5 μl of 500 μM L-DOPA. Next, 90 μl of ABTS solution

was added to the well, and the absorbance was read immediately at 734 nm. Synergy was

calculated from this data using the following formula:

$$\text{Synergy Ratio} = \frac{(\Delta \text{Abs 734 Compound Alone} + \Delta \text{Abs 734 DOPA Alone})}{\Delta \text{Abs 734 Compound with DOPA}}$$

Glyphosate Effect on L-DOPA

To determine if L-DOPA is reacting with glyphosate, we analyzed by NMR. We diluted

300 mM stock of glyphosate in water to 60 mM (10 mg/ml) in D_2O, prepared 20 mM (4 mg/ml) L-

DOPA in D_2O, and prepared two mixtures of glyphosate and L-DOPA: one with 20 mM (4

mg/ml) L-DOPA and 60 mM (10 mg/ml) of glyphosate in D_2O, and another with a low

concentration of 1 mg/ml for both compounds equaling 5 mM L-DOPA and 6 mM glyphosate.

We then performed ^{31}P-NMR and ^{1}H-NMR on these samples.

Glyphosate Effect on Tyrosinase

To determine the tyrosinase kinetics with glyphosate as an inhibitor, we serially diluted

155 μl of 20 mM L-DOPA in Milli-Q water. To each dilution of L-DOPA we added 20 μl of

glyphosate diluted in PBS and 5 μl of 2 μg/ml mushroom tyrosinase to the reaction mix. In order

to account for non-enzymatic oxidation of L-DOPA, we ran an experiment in parallel, in which

we added 5 μl of Milli-Q water instead of tyrosinase. The reaction mix was kept at 30°C for 24 h.

The plate was read at 490 nm. To calculate enzyme-specific oxidation of L-DOPA, the no

enzyme values were subtracted from the tyrosinase rows. The kinetics curve is plotted as a

function of absorbance after 24 h of reaction time versus concentration of L-DOPA.

We tested if tyrosinase concentration had an effect on the percent inhibition of the

reaction. We prepared dilutions of tyrosinase. We added 5 μl of each dilution to a 96-well plate,

and added 135 μl of Milli-Q water, 20 μl of 20 mM L-DOPA, and 20 μl of glyphosate in PBS. We

measured maximum velocity of this reaction at 490 nm. The difference in velocities and percent

inhibition reported were calculated by difference = $V_{max\ water}$ - $V_{max\ glyph}$, and percent inhibition =

$100*(V_{max\ glyph}/V_{max\ water})$.

To determine if glyphosate irreversibly affects tyrosinase activity, 450 µl of 20 µg/ml

mushroom tyrosinase was prepared in 450 µl of 50 mM sodium phosphate buffer, pH 7, either

with 50 µl of 300 mM glyphosate, or 50 µl of Milli-Q water. The enzyme solution was loaded into

a hydrated 10,000 MWCO Slide-a-lyzer dialysis cassette (Thermo Scientific), and the enzyme

solutions were dialyzed in a 50 mM sodium phosphate buffer at 4°C, according to the

manufacturer's protocol. Protein concentrations were measured and normalized using sodium

phosphate buffer. To measure the kinetics of the control enzyme versus the treated enzyme, a

kinetics assay was performed as previously described. Each reaction's maximum velocity is

determined and plotted.

Copper Rescue of Melanin Inhibition

As previously described, serial dilutions of glyphosate were arrayed in eight rows; one

row per copper ion concentrations to be tested. Copper sulfate was prepared and serially diluted

and 10 µL of the copper solution is added to each well containing the glyphosate dilution. To

each well 150 µL of reaction mix (125 µL of Milli-Q water, 20 µL of 20 mM L-DOPA, and 5 µL of

2 µg/mL mushroom tyrosinase (5 µl of water used for auto-oxidation experiments) was added.

The final copper ion concentrations were 400, 200, 100, 50, 25, 12.5, 6.25, and 0 µM. The

dopachrome and melanin measurements are reported as previously described.

Cyclic Voltammetry

Cyclic voltammetry was performed using a Metrohm Autolab potentiostat (Switzerland),

3 mm Glassy Carbon working electrode, 10 mm x 10 mm x 0.1 mm platinum plate counter

electrode, and an Ag/AgCl reference electrode in 3 M KCl solution. Solutions were prepared in

0.1x PBS (Difco) at a pH 6.00, adjusted with NaOH and HCl. 10 mL of L-DOPA solution was

freshly prepared in this buffer, and 1 mL of glyphosate, glycine, water, etc, solution at pH 6.00

were added to the L-DOPA. Readings were done with three tracings at a scan rate of 50 mV/s

at intervals of 5 mV steps. Glassy carbon electrode was washed and polished between readings

with slurry of alumina powder and water on cloth pads.

DATA AVAILABILITY

The 16S rRNA sequencing datasets generated during this study are available at Mendeley Data

at DOI:10.17632/6ymh76hmzm.1, and the datasets from the remaining experiments are

available at Mendeley Data at DOI: 10.17632/xndcmbn6wd.2.

FIGURE LEGENDS

Fig. 1. *Glyphosate Inhibits G. mellonella Melanization and Increases Infection*

***Susceptibility.* (A**) Glyphosate inhibits the phenoloxidase activity of 1:10 dilutions of

hemolymph without exogenously added L-DOPA. (**B**) AMPA, a primary metabolite of

glyphosate, inhibits *G. mellonella* phenoloxidase-mediated melanization similar to glyphosate.

Error bars in **(A-B)** represent ± SD (**C**) *G. mellonella* larvae drugged with glyphosate solution

(10 µl of 1 mM) in PBS and infected 5 h post treatment with 10^4 cells of WT *C. neoformans* die

rapidly compared to PBS-treated controls. Death events were recorded daily. Each infection

condition represents survival of 95 animals, pooled together from four biological replicates, and

six total technical replicates. Statistical significance was assessed by Gehan-Breslow-Wilcoxon

test, which we used to place weight on early timepoints in the survival curve. We used this test

because we expected to see the glyphosate-mediated differences early in the infection due to

the timing of the glyphosate treatment. Since w where the expected effects of the one-time pre-

treatment with glyphosate would be. **(D)** The size of the dark melanized particles within nodules

upon *C. neoformans* infection are significantly smaller in the glyphosate-treated (10 µl of 1 mM)

infected groups compared to the PBS-treated infected groups, which were analyzed for

significance using a nested non-parametric Mann-Whitney-Wilcoxon rank test. Horizontal bar

represents the median value and the error bars represent the 95% Confidence Interval **(E)** The

degree of melanin encapsulation of the yeast within the nodule is also reduced in the

glyphosate-treated (10 µl of 1 mM) groups, as measured on a scale of 0 (no melanin

encapsulation) to 4 (very high levels of melanin encapsulation) as demonstrated in (**F**).

Numbers in each bar represent the number of encapsulated *C. neoformans* for each score.

Statistical significance was assessed using a Chi-squared table test. Data in (**D**) and (**E**)

represent data over three independent replicates with three larvae used per condition per

replicate. (**G**) Representative brightfield micrographs showing the hemocyte and nodule

formation in the different treatment groups at 20x and 100x magnification. Scale bars represent

10 µm. Nested non-parametric Mann-Whitney-Wilcoxon rank test performed using R for R 4.0.2

GUI 1.72 for Mac OS at https://www.r-project.org/ (R Core Team, 2020) and the *nestedRanksTest*

package (*Version 0.2*, D.G. Scofield , 2014)(134). All other statistical analyses performed using

GraphPad Prism version 8.4.3 for Mac OS, GraphPad Software, San Diego California USA,

www.graphpad.com. See also, Supplementary Fig. 1.

Fig. 2. *Glyphosate Effects on A. gambiae Phenoloxidase Activity and Susceptibility to*

Plasmodium Infection. (**A**) Glyphosate inhibits phenoloxidase activity in *A. gambiae* homogenate.

Enzyme activity represents three biological replicates with three technical replicates for each

condition. (**B)** Glyphosate treatment increases the susceptibility of the *A. gambiae* to *P. falciparum*

infection as measured by oocyst count per midgut. Increased glyphosate doses are associated with

increased median oocyst burden. Parasite infection represents four biological replicates and four

separate infections, line indicates median, and differences in parasite burden analyzed for

significance using non-parametric Kruskal–Wallis test with each group compared to the control

group with Dunn's correction for multiple comparisons. (**C**) Low doses of glyphosate enhance the

survival of adult mosquitoes, while the higher doses diminish their survival as compared to the

control. Survival curves represent 120 animals from three independent replicates composed of

groups of 40 mosquitoes, and survival was examined for statistical significance using the Log-Rank

Mantel-Cox analysis with a Bonferoni correction for multiple comparisons. (D) Hazard ratios

calculated from the Cox Mixed Effects Model to account for fixed (glyphosate treatment) and

random effects (replicate). Hazard ratios<1 indicate lower risk of death compared to control values,

and values >1 indicate a higher risk of death compared to the control. Hazard Ratio of 1 is depicted

by a dotted line. The Cox Mixed Effects modeling was performed using R for R 4.0.2 GUI 1.72 for

Mac OS at https://www.r-project.org/ (R Core Team, 2020) and the *coxme* package (*Version 2.2-16*,

T.M. Therneau, 2020)(141). All other statistical analyses performed using GraphPad Prism

version 8.4.3 for Mac OS, GraphPad Software, San Diego California USA, www.graphpad.com.

The statistical significance in **(C)** is coded as: ns - p > 0.05, * - p<0.05, ** - p<0.01, *** -p<0.001,

and **** - p<0.0001. See also Supplementary Fig. 2.

Fig. 3. *Glyphosate Alters the Composition, but Not Density, of the A. gambiae Midgut*

Microbiota. (**A**) Glyphosate does not alter microbial density of the culturable mosquito midgut

bacteria (grown on LB agar). Each sample consists of 40-50 individual mosquito midguts over three

independent replicates. Error bars represent the mean and ±SD. **(B)** Glyphosate alters the

composition of the mosquito microbiota, leading to decrease of Entereobacteriacae and an increase

of *Asaia* spp. **(C)** The glyphosate treatments do not significantly alter alpha diversity as measured by

the Shannon Index (statistical analysis conducted using one-way ANOVA; NS = p >0.05). **(D)**

However, the glyphosate-treated and control-treated microbiota form distinct clusters in principle

coordinates analysis, measured by Bray-Curtis dissimilarity. Statistical significance was tested by

PERMANOVA (p <0.001, R=0.557). Each treatment group represents 5 individual mosquito

midguts. For more information see also Supplementary Fig. 3.

Fig. 4. *Glyphosate Inhibits in vitro Melanin Production.* (A) An overall schematic of the

Mason-Raper pathway of melanization mediated by tyrosinase and auto-oxidation. (**B**)

Glyphosate inhibits formation of dopaquinone produced by tyrosinase-mediated and

spontaneous oxidation of L-DOPA. Dopaquinone is indicated by the absorbance of an MBTH-

Dopaquinone adduct pigment at 505 nm. Absorption levels are shown relative to the no

glyphosate control with background (MBTH mixture) subtracted after 1 h at 30ºC **(C)** Glyphosate

decreases the rate of dopachrome formation and inhibits dopachrome production from

tyrosinase oxidation of L-DOPA. Rate of dopachrome formation is the reaction V_{max} at 490 nm

relative to the V_{max} without glyphosate. Dopachrome production is shown as the absorbance at

490 nm relative to the control after 30 min of reaction. **(D)** Melanin production is inhibited by

glyphosate with tyrosinase and auto-oxidation of L-DOPA. Melanin levels are measured as the

absorbance at 490 nm after 5 d of reaction. Inset shows a representative image of the data in

(**D**) showing melanization inhibition with increasing glyphosate concentration. Values are

depicted relative to the no glyphosate control. Error bars represent ±SD. Each experiment was

performed at least three independent replicates.

Fig. 5. *Phosphate-Containing Compounds Inhibited Melanization Similarly to Glyphosate.*

Glyphosate, *o*-phosphoserine (PS), phosphonoacetic acid (PAA), pyrophosphate (pyro), and

phosphoric acid (PA) inhibit dopaquinone formation **(A)**, rate of dopachrome formation **(B)** and

dopachrome levels **(C)**, and melanin formation **(D)**, whereas their respective non-phosphate

analogs, glycine (gly), serine (ser), and acetic acid (AA) do not inhibit any step of melanization

(A-D). (**E**) Auto-oxidation of L-DOPA is inhibited by glyphosate, PS, PAA, Pyro, and PA in a

similar manner. The compounds tested (**F)** were diluted in 300 mM stock solution and titrated to

pH between 5 and 6. Absorption and rates are shown relative to the internal no drug control.

Grayscale bars represent mean absorbance at 490 nm relative to no compound control. The

darker colors correspond to increased pigment formation. Inset shows a representative image of

the data in (**E**) showing the effects of the compounds on auto-oxidation. Error bars in (**A-C)**

represent ±SD. Each experiment represents at least three independent replicates.

Fig. 6. *Glyphosate Does Not Directly Inhibit Tyrosinase Activity* (A). Tyrosinase activity is

not irreversibly inhibited and glyphosate-treated enzyme has normal activity when glyphosate is

dialyzed out of solution. **(B)**. Glyphosate appears as a non-competitive inhibitor of tyrosinase in

Michaelis-Menten kinetics assays measuring the change in absorbance at 490 nm over 24 h

compared to the no tyrosinase background. **(C)** The rate of dopachrome formation with

glyphosate treatment is smaller than the slope of the control treatment across all concentrations

of tyrosinase. This reduced slope indicates reversible inhibition. The assay is performed under

constant L-DOPA and glyphosate concentrations. Shaded areas represent the 95% CI of the

linear regression **(D)** Adding Cu^{+2} to L-DOPA-tyrosinase reactions with glyphosate does not

rescue melanin inhibition compared to the glyphosate-free control. (See also Supplementary Fig.

5) Grayscale bars represent mean absorbance at 490 nm relative to no glyphosate and no

copper control. The darker colors correspond to increased pigment formation. Error bars in **(A-**

C) represent ±SD. Each experiment represents at least three independent replicates.

Fig. 7. *Glyphosate Affects the Oxidative Properties of Melanogenesis.* (A) None of the

melanin inhibitors exhibit radical quenching properties in an ABTS assay aside from glyphosate,

which shows weak antioxidant properties after several hours in the ABTS solution. Absorbance at

734 nm is an indicator of how much ABTS remains in radical form (not quenched). **(B-C)** Citric acid

(CA), a non-radical quenching antioxidant (antioxidant synergist) exhibits similar melanin inhibition

as glyphosate and phosphoric acid, another known antioxidant synergist. Grayscale bars in **(C)**

represent absorbance at 490 nm relative to no compound control, with the darker colors

correspond to increased pigment formation. **(D)** Glyphosate, phosphoserine, phosphoric acid, and

citric acid show synergy with the antioxidant L-DOPA. The addition of these compounds to L-DOPA

enhances its radical quenching abilities by approximately 50%. Black dotted line represents the

normalized ABTS absorbance treated with water. The other compounds tested here alone do not

show much deviation from this line. The blue dotted line indicates the ABTS solution treated with L-

DOPA alone. ABTS treated with L-DOPA and synergetic compounds together are below this line. (**E**)

Average cyclic voltammogram showing the changes in oxidation and reduction of L-DOPA and

dopaquinone when exposed to 16 mM glyphosate but not water. Numbers correspond to shifted

peaks or peaks with less current compared to the water control. Peak 1 corresponds to L-DOPA

oxidation (**F**); Peak 2 likely corresponds to dopaquinone reduction (**G**). glyphosate shifts Peak 1 and

2 toward a decreased redox potential and diminishes the current of Peak 1 and 2 in a dose-

dependent manner (**H**) - notably decreasing Peak 2 current intensity to the point of non-existence

(**I**). Each experiment represents at least three independent replicates. Error bars in (**A-B, D**)

represent ±SD. See also Supplementary Fig. 6 and 8 .

Supplementary Fig. 1. *G. mellonella* **Supplemental Data. (A)** Broad-spectrum protease

inhibitor (cOmplete, Roche) was added to *G. mellonella* hemolymph to prevent the activation of

new phenoloxidase, and to control for any impact that glyphosate may have on phenoloxidase

activation cascade, cell viability, and gene expression. The general trend remains the same that

glyphosate inhibits phenoloxidase activity with and without protease inhibitor, albeit lower with

protease inhibitor due to the lower concentration of activated enzyme. **(B)** Phenoloxidase

activity was assessed using exogenous L-DOPA for one batch of *G. mellonella,* during these

experiments, the lower concentration of glyphosate resulted in increased phenoloxidase activity

as compared to the control. This suggests that there may be some cellular regulation of

phenoloxidase induced by glyphosate. It is possible that the doses of glyphosate tested elicit

some cellular response that increases phenoloxidase expression, secretion, and/or activation as

a feedback/hormesis-like response to the reduced melanin production. These data represent

three independent replicates, but this pattern of enzymatic activity as a function of glyphosate

concentration was not seen in subsequent batches of larvae. (**C**) Hemocyte viability was not

dramatically affected by concentrations of glyphosate ranging from 100 μM to 10 mM, indicating

that our data are likely not artifacts of cytotoxic concentrations of glyphosate. Error bars in **(A-C)**

represent ± SD. (**D)** AMPA, a major metabolite of glyphosate, inhibits tyrosinase-mediated

melanization similar to glyphosate. Grayscale bars represent mean absorbance at 490 nm

relative to no compound control. The darker colors correspond to increased pigment formation.

(E) Larvae treated with glyphosate and subsequently infected with *lac1Δ* mutant *C. neoformans*

strain showed a similar pattern of increased susceptibility as the wild type H99, although the

differences in susceptibility with the *lac1Δ* infected larvae are not statistically significant. Each

experiment represents at least three independent replicates. The PBS mock infection condition

represents survival of 95 animals, over the span of four biological replicates, and six total

technical replicates. The *lac1Δ* mutant infection represents survival of 75 animals over the span

of four biological replicates. The PBS mock infection data is the same as the data in Fig. 5b, as

all the infections were done concurrently under the same conditions. (**F)** Single injection of 10 μl

of 1mM glyphosate does not affect the pupation of *G. mellonella* at 30°C and room temperature

(RT). Data from 30°C represents 25-35 animals for each group over two biological replicates,

and data from RT represents 45 animals from each group over three biological replicates.

Statistical analysis performed using Log-rank Mantel-Cox tests. (**G-I**) The three individual

replicates from **Fig. 1D** showing the size of the dark melanized particles within nodules are

significantly smaller in the glyphosate-treated infected groups compared to the PBS-treated

infected groups, with (**G**) and **(H)** showing that there were more melanized spots in the PBS-

treated infected group compared to the glyphosate-treated. All statistical analyses performed

using GraphPad Prism version 8.4.3 for Mac OS, GraphPad Software, San Diego California

USA, www.graphpad.com.

Supplementary Fig. 2 Low efficiency *Plasmodium falciparum* infection of *A. gambiae and*

effects on mosquito cuticle (**A)**. Oocyst count per midgut for mosquitoes treated with or

without glyphosate and infected with high-passage *Plasmodium falciparum* gametocyte culture,

resulting in a low efficiency infection. Data represents one biological replicate. Dotted black line

indicates y=0. Black lines for each condition indicate median oocyst count per midgut. We have

chosen not to include the data from this replicate in the data shown in Fig. 6, because the

results from this one-off replicate appear due to poorly infectious parasite culture. Additionally, it

is difficult to make comparisons using the low infection burden of the control group with a with

the treatment groups, as well other replicates with higher oocyst burdens. **(B)** Infection

prevalence (percent midguts with at least one oocyst) from the experiment described in (**A**).

Fisher's Exact test performed for each condition individually compared to control and corrected

for multiple comparisons using the Bonferroni method. **(C)** 5 days of 1 mM glyphosate treatment

in adult female mosquitoes does not influence the abdomen's cuticular darkness as measured

by mean gray value with 0 being pure black and 255 being pure white. Data representative of 2

biological replicates with 88 mosquitoes measured per condition. **(D)** Wing length, as a proxy for

body mass and size, is not affected by 5 days of glyphosate treatment. Data representative of a

single biological replicates with 32-36 mosquitoes measured per condition. Line and error bar

represent mean ± SD in **(C-D).** Unpaired t-test performed to determine statistical significance in

(C-D). All statistical analyses performed using GraphPad Prism version 8.4.3 for Mac OS,

GraphPad Software, San Diego California USA, www.graphpad.com.

1013

Supplementary Fig. 3. *Glyphosate affects the A. gambiae microbiota in a dose-*

independent manner **(A)** At the class level, glyphosate leads to an enrichment of

Alphaproteobacteria and a depletion in Gammaproteobacteria**.** Tables showing the relative

abundance of bacterial classes (**B**) and individual bacterial genera (**C**) following glyphosate

treatment . (**D)** Alpha diversity does not follow a distinctive pattern with increasing glyphosate

dose. **(E)** Glyphosate-treated and control-treated microbiota cluster separately in ordination

space, but the clusters are not dose-dependent.

1021

1022   **Supplementary Fig 4.** *Reaction of glyphosate with L-DOPA.* Representative ^{1}H NMR spectra

1023   of 60 mM glyphosate solution in D_2O (**Green**), 20 mM L-DOPA solution in D_2O (**Red)**, and 20

1024   mM L-DOPA mixed with 60 mM glyphosate in D_2O (**Blue**). There appears to be no shift in ^{1}H

1025   peaks and no appearance of new peaks, which is indicative of no reaction occurring between

1026   the compounds. Data representative of three independent replicates

1027

1028   **Supplementary Fig. 5.** *Glyphosate appears to "buffer" copper concentration in solution.*

1029   High doses (2-16 mM) of glyphosate prevent the enzymatic activity enhancing effects of lower

1030   copper concentration (6.25-25 µM), but high doses of glyphosate also prevent the enzyme

1031   inhibitory effects of high copper concentration (100-400 µM)**.** Error bars represent ±SD. Data

1032   represents at least three independent replicates.

1033

1034   **Supplementary Fig. 6.** *Antioxidant Properties of glyphosate.* (**A**) Change in absorbance of

1035   ABTS solution at 734 nm over time for 33.33 mM glyphosate relative to the no glyphosate

1036   control. This indicates glyphosate quenches free radicals over an extended period of time. (**B**)

1037   Calculated antioxidant radical scavenging synergy between compounds tested and L-DOPA.

1038   Values represent the mean of at least three independent replicates. Error bars represent ±SD.

1039

Supplementary Fig. 7. *Glyphosate inhibits melanin production independent of L-DOPA*

concentration. (**A**) Inhibitory concentrations of glyphosate are not affected by L-DOPA

concentration. This indicates that glyphosate is not reacting proportionately with L-DOPA as

measured by absorbance at 490 nm after 5 d of reaction, relative to the no glyphosate control

and with background absorbance subtracted. (**B**) The IC_{50} of glyphosate remains constant at

approximately 1 mM relative inhibition of melanin production appears dependent on glyphosate

concentration alone, and not on L-DOPA to glyphosate ratio. Error bars represent ±SD. Each

experiment represents at least three independent replicates. Grayscale bars represent mean

absorbance at 490 nm relative to no compound control. The darker colors correspond to

increased pigment formation. Red line represents the approximate IC_{50}. Crossed out boxes

represent values with no data.

1051

Supplementary Fig. 8. Cyclic Voltammetry Supplemental Data. (A) Peak 1 was validated as

the oxidation of L-DOPA, and Peak 2 was validated as the reduction peak of dopaquinone by

correlating increased peak intensity with increasing concentration of L-DOPA under the same

potentiostat parameters. **(B)** Glycine (16 mM) – a non-phosphate analog of glyphosate, a non-

inhibitor of melanization, and a non-antioxidant - does not alter the oxidation potential of L-

DOPA. Conversely, Citric Acid (16 mM) – a known synergistic antioxidant and inhibitor of

melanization – does alter the oxidation potential of L-DOPA in similar ways as glyphosate. Each

experiment represents at least three independent replicates, with three cycles per replicate. The

tracings represent the mean value of the three replicates over the course of three cycles.

1061

Supplementary Fig. 9. Experimental Methods Diagram for Insect Experiments. (A). During *G.*

mellonella infection with *C. neoformans,* larvae were injected with 10 µl of 1 mM glyphosate, left to

recover for 5 h, and were subsequently infected with 10^4 cells/larvae of *C. neoformans* H99 strain.

Survival was monitored for 14 days. (**B**) During *A. gambiae* infection with *P. falciparum,* mosquitoes

were drugged with glyphosate-laced 10% sucrose solution for 5 days, then fed with a *P. falciparum-*

infected blood meal, and fed 10% sucrose for 8 d. On Day 8, mosquitoes were dissected, and the

midguts were stained with mercurochrome to facilitate oocyst enumeration. **(C)** glyphosate-drugged

mosquitoes were dissected under sterile conditions, and five midguts were collected individually per

condition. DNA was extracted from samples and bacterial 16S rRNA genes were amplified by PCR

and sample-specific Illumina adapters were ligated to products. PCR products were pooled and

sequenced on the Illumina MiSeq platform. Data were then analyzed using mothur to construct

contigs, align reads, remove ambiguous bases and chimeric regions, align sequences to the Silva

16S V4 reference database, and cluster reads into 3% operational taxonomic units (OTUs).

Sequences from known contaminants were removed. Alpha and beta diversity measurements were

performed using the Shannon diversity index and Bray-Curtis dissimilarity distance respectively and

plotted using MicrobiomeAnalyst. Figures made with BioRender.

1078

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Figure 1

A. *G. mellonella* PO Activity

Relative V_{max} (490 nm) vs Glyphosate (mM)

n = 7

B. *G. mellonella* PO Activity

Relative V_{max} (490 nm) vs Glyphosate (mM)

- Glyph
- AMPA
- Water

C. *G. mellonella* Infection with *C. neoformans*

Percent survival vs Days

- PBS + PBS
- Glyph + PBS } ns
- PBS + H99
- Glyph + H99 } p= 0.013

n = 95/group

D. Melanized Particle Area

Melanized Particle Area (pixels^2)

p = 0.004

PBS   H99-PBS   H99-Glyph

Melanization Score

Fungal Cells within Nodules

H99-PBS: 117, 39, 60, 58, 33
H99-Glyph: 170, 43, 81, 74, 14

- 4
- 3
- 2
- 1
- 0

Representative Scoring References

10 µm

G.

	PBS	H99-PBS	H99-Glyph
20x			
100x			

Figure 2

Figure 3

Figure 4

A

L-DOPA →(**Auto-oxidation** / Tyrosinase)→ Dopaquinone (DQ) → Cyclodopa

Cyclodopa →(Radical-mediated Redox Exchange; DQ → L-DOPA)→ Dopachrome (DC) → Dihydroxyindole →(ROS)→ Eumelanin

B — Dopaquinone Production
- Absorbance (505 nm) vs Glyphosate (mM)
- Tyrosinase
- Auto-oxidation

C — Dopachrome Formation
- Relative Absorbance (490 nm) / Relative V_{max} (490 nm) vs Glyphosate (mM)
- Level
- Rate

D — Melanin Production
- Relative Absorbance (490 nm) vs Glyphosate (mM)
- Glyphosate Concentration
- Tyrosinase
- Auto-oxidation

Figure 5

Figure 6

Figure 7

A Radical Scavenging Properties

B Dopachrome Production

C Melanin Production

D Acid Synergism With L-DOPA

E L-DOPA Voltammograms

F Peak 1: L-DOPA Oxidation

L-DOPA → Dopaquinone

G Peak 2: Dopaquinone Reduction

Dopaquinone → L-DOPA

H Dose Dependent Effect of Glyphosate

I

Supplementary Figure 1

Supplementary Figure 2

A. Low *P. falciparum* Infection Burden

B. Low *P. falciparum* Infection Prevalence

C. Mosquito Cuticle Pigmentation

D. Mosquito Wing Length

Supplementary Figure 3

A

D

B

	[Glyphosate]			
Class	- Glyphosate	30 µM	300 µM	3 mM
Actinobacteria	0.0058	0.005642	0.002504	0.00476
Alphaproteobacteria	0.45129705	0.6918657	0.7944112	0.5505296
Bacilli	0.01333963	0.0286482	0.006664	0.0113333
Betaproteobacteria	0.0041215	0.000844	0.0001336	0.000471
Gammaproteobacteria	0.51917302	0.2566697	0.1942569	0.423504
Negativicutes	0.00042991	0.0025054	0.0003698	0.0005008
Unclassified Proteobacteria	0.00580703	0.013826	0.00166	0.0088853

C

	[Glyphosate]			
Genus	- Glyphosate	30 µM	300 µM	3 mM
Asaia	0.1336578	0.664002	0.789541	0.442692
Unclassified Enterobacteriaceae	0.58495882	0.155066	0.174655	0.130596
Pseudomonas	0.02301406	0.063686	0.01635	0.271084
Unclassified Acetobacteraceae	0.21889554	0.006761	0.000116	0.002701
Unclassified Sphingomonadaceae	0	0	0.001265	0.087311
Acetobacter	0.00847871	0.021091	0.003683	0.01762
Unclassified Gammaproteobacteria	0.00343976	0.028129	0.000996	0.00285
Unclassified Proteobacteria	0.00464562	0.013806	0.001663	0.008885
Streptococcus	0.00243342	0.013	0.002609	0.003708
Unclassified Bacillales	0.0039732	0.007579	0.00253	0.004348
Others	0.0164868	0.026856	0.006594	0.02819

E

Supplementary Figure 4

Supplementary Figure 5

Glyphosate "Buffers" Copper Ions

Glyphosate Dose
- 16mM
- 4mM
- 2mM
- 1mM
- 0.5mM
- 0mM

V_{max} (490 nm)

Concentration Cu^{+2} (μM)

Supplementary Figure 6

A

Glyphosate Effect on ABTS Absorbance

B

Compound	% Radical Quench-Compound Alone	% Radical Quench-Compound and L-DOPA	Synergy (Compared with L-DOPA alone)
Phosphoacetic Acid	0.020	72.288	0.483
Citric Acid	1.755	72.320	0.507
Phosphoric Acid	1.096	70.831	0.508
Pyrophosphate	3.320	72.675	0.526
Acetic Acid	1.327	64.155	0.565
Glyphosate	5.839	69.483	0.586
Phosphoserine	1.558	57.443	0.635
Glycine	-0.566	44.252	0.776
Serine	-0.386	40.808	0.846
Water	0.000	36.635	0.953
L-DOPA		34.904	1.000

Supplementary Figure 7

Supplementary Figure 8

A   L-DOPA Peak Validation

- 0 mM
- 1 mM
- 2.5 mM
- 5 mM
- 10 mM

Peak 1

Peak 2

Current (µA)

Potential applied (V)

B   Positive and Negative Inhibitor Controls

- L-DOPA
- Glyphosate
- Glycine
- Citric Acid

Current (µA)

Potential applied (V)

Supplementary Figure 9